

SEQUENCE LISTING

```
<110> Agarwal, Pankaj
      Lee, Judithann M.
      Smith, Randall F.
      White, John R.
<120> NOVEL COMPOUNDS
<130> GP50029-1
<140> 10/687,268
<141> 2003-10-15
<150> 60/213,161
<151> 2000-06-22
<150> 60/213,156
<151> 2000-06-22
<150> PCT/US01/19929
<151> 2001-06-22
<150> 10/312,088
<151> 2002-12-20
<160> 46
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 1383
<212> DNA
<213> Homo sapiens
<400> 1
atgcttggaa tttggattgt tgcattcttg ttctttggca catcaagagg aaaagaagtt 60
tgctatgaaa ggttagggtg tttcaaagat ggtttaccat ggaccaggac tttctcaaca 120
gagttggtag gtttaccctg gtctccagag aagataaaca ctcgtttcct gctctacact 180
atacacaate ccaatgeeta teaggagate agtgeggtta attetteaae tatecaagee 240
tcatattttg gaacagacaa gatcacccgt atcaacatag ctggatggaa aacagatggc 300
aaatggcaga gagacatgtg caatgtgttg ctacagctgg aagatataaa ttgcattaat 360
ttagattgga tcaacggttc acgggaatac atccatgctg taaacaatct ccqtqttqtt 420
ggtgctgagg tggcttattt tattgatgtt ctcatgaaaa aatttgaata ttccccttct 480
aaagtgcact tgattggcca cagcttggga gcacacctgg ctggggaagc tgggtcaagg 540
ataccaggcc ttggaagaat aactgggttg gacccagctg ggccattttt ccacaacact 600
ccaaaggaag tcaggctaga cccctcggat gccaactttg ttgacgttat tcatacaaat 660
gcagctcgca tcctctttga gcttggtgtt ggaaccattg atgcttgtgg tcatcttgac 720
ttttacccaa atggagggaa gcacatgcca ggatgtgaag acttaattac acctttactg 780
aaatttaact tcaatgctta caaaaaagaa atggcttcct tctttgactg taaccatgcc 840
cgaagttatc aattttatgc tgaaagcatt cttaatcctg atgcatttat tgcttatcct 900
tgtagatcct acacatcttt taaaqcaqqt acatqtqtaq qatqtqcaqa tttqttacat 960
aggatagata agataggaag tcatacttcc catgtgtttt taaccctttc tctccctttc 1020
```

cttcttgttt ccttatatct aggttggagg cacaaattgt ctgttaaact cagtggaagc 1080

```
gaagtcactc aaggaactgt ctttcttcgt gtaggcgggg cagttaggaa aactggggag 1140
tttgccattg tcagtggaaa acttgagcca ggcatgactt acacaaaatt aatcgatgca 1200
gatgttaacg ttggaaacat tacaagtgtt cagttcatct ggaaaaaaca tttgtttgaa 1260
gattctcaga ataagttggg agcagaaatg gtgataaata catctgggaa atatggatat 1320
aaatctacct tctgtagcca agacattatg ggacctaata ttctccagaa cctgaaacca 1380
                                                                  1383
tgc
<210> 2
<211> 927
<212> DNA
<213> Homo sapiens
<400> 2
atgccgttcc tgcagctgaa agggagagca acacctccat cctggagaca cgatagccgc 60
tcacttgttc acctgctgga cggcaaggag ggcgtgtggg acaccacggg ctatgcctta 120
gggagcagag aatcattgaa tcctgacatg gggattggtg acccacatgg acacagcact 180
gtccacacca gggaagcagg cactgcctgt ccattacagc ttctaggtgc tcgggaggcc 240
agtotgotgg cotgtgggat otgocaggoo totggccaaa tottcatcac ocaaaccotg 300
gggatcaagg gatatcggac tgtcgtggcc ctggataagg tccctgagga tgttcaggaa 360
tacagctggt actggggtgc aaacgacagc gcaggaaaca tgattatcag ccacaaaccg 420
cccagtgccc agcagcctgg gcccatgtac actggcaggg agagagtgaa cagagaaggc 480
agcctgttga tcaggccgac tgcattaaat gacacgggaa actacactgt tcgggtggtt 540
gcaggcaatg agacccaaag agcaaccggc tggctggagg ttctagatgg gcccgactat 600
gtgctgctga ggagcaatcc tgatgatttc aacggcattg tgacagctga gatcggctcc 660
caagtggaaa tggagtgtat ctgctattcc ttcctggatc tcaagtacca ctggatccac 720
aatggctccc tcctgaactt ctcagatgca aagatgaacc tctcgagtct tgcctgggag 780
cagatgggcc gttaccgatg cactgtggag aaccccgtga cacagctgat catgtacatg 840
gacgtcagga tccaggcccc ccatgagtgc agcagctccc ctccaggctc atgctttgca 900
catctccctg cctccatgcc ctgctag
<210> 3
<211> 1374
<212> DNA
<213> Homo sapiens
<400> 3
atggacettt ccagacecag atggagectg tggaggaggg tetteeteat ggecagtetg 60
ctggcctgtg ggatctgcca ggcctctggc caaatcttca tcacccaaac cctggggatc 120
aagggatate ggactgtegt ggeeetggat aaggteeetg aggatgttea ggaataeage 180
tggtactggg gtgcaaacga cagcgcagga aacatgatta tcagccacaa accgcccagt 240
gcccagcage etgggcccat gtacactggc agggagagag tgaacagaga aggcagcctg 300
ttgatcaggc cgactgcatt aaatgacacg ggaaactaca ctgttcgggt ggttgcaggc 360
aatgagaccc aaagagcaac cggctggctg gaggttctag agttgggaag caatctgggc 420
atctccgtca atgccagctc cctggtggag aacatggatt ctgtggctgc tgactgcctc 480
acaaatgtca ccaacatcac gtggtatgtg aatgatgtgc ctacctctag tagtgaccgg 540
atgacaattt ccccagacgg caagacctc gtcatcctca gggtcagccg ctatgacaga 600
acaattcagt gcatgataga gagtttccca gagatctttc agagaagtga acgcatctct 660
ctgactgtgg cctatgggcc cgactatgtg ctgctgagga gcaatcctga tgatttcaac 720
ggcattgtga cagctgagat cggctcccaa gtggaaatgg agtgtatctg ctattccttc 780
ctggatctca agtaccactg gatccacaat ggctccctcc tgaacttctc agatgcaaag 840
atgaacctct cgagtcttgc ctgggagcag atgggccgtt accgatgcac tgtggagaac 900
cccgtgacac agctgatcat gtacatggac gtcaggatcc aggcccccca tgagtgtcct 960
cttccttcag ggatcttacc tgttgtccac agagatttct ccatctcagg atccatggtg 1020
atgttcctca tcatgctgac agtgctgggt ggcqtttaca tctgtggagt cctgatccat 1080
gctctgatca accactactc aatcaggtgc cctcattgct ctgggacaag ggtgggatgt 1140
tggctggggg ctgggactca ggagccagcc ctccctccag aggggaagca gagccagaaq 1200
gggagggata agccaggaac taggttgtca gggatcatct ggggcagaca gatcagcccc 1260
```

```
caggacctga agctgatggg agcaagagag ggtttagagt cggccatggt tctaaatagc 1320
tgtggggttt cttctagcaa cttcccttct ctttgtgttt ataagggata ttaa
<210> 4
<211> 2115
<212> DNA
<213> Homo sapiens
<400> 4
atgctccatg atgggttgac tgcacctgat gggtgtggaa tctacagcct gaccgggcgg 60
gaagtcctga cgcccttccc aggattgggc actgcggcag ccccggcaca gggcggggcc 120
cacctgaagc agtgtgacct gctgaagctg tcccggcggc agaagcagct ctgccggagg 180
gageceggee tggetgagae cetgagggat getgegeace teggeetget tgagtgeeag 240
tttcagttcc ggcatgagcg ctggaactgt agcctggagg gcaggatggg cctgctcaag 300
agaggettea aagagacage ttteetgtae geggtgteet etgeegeeet eacceaeae 360
ctggcccggg cctgcagcgc tgggcgcatg gagcgctgca cctgtgatga ctctccgggg 420
ctggagagcc ggcaggcctg gcagtggggc gtgtgcggtg acaacctcaa gtacagcacc 480
aagtttctga gcaacttcct ggggtccaag agaggaaaca aggacctgcg ggcacgggca 540
gacgcccaca atacccacgt gggcatcaag gctgtgaaga gtggcctcag gaccacgtgt 600
aagtgccatg gcgtatcagg ctcctgtgcc gtgcgcacct gctggaagca gctctccccg 660
ttccgtgaga cgggccaggt gctgaaactg cgctatgact cggctgtcaa ggtgtccagt 720
gccaccaatg aggccttggg ccgcctagag ctgtgggccc ctgccaggca gggcagcctc 780
accaaaggcc tggccccaag gtctggggac ctggtgtaca tggaggactc acccagcttc 840
tgccggccca gcaagtactc acctggcaca gcaggtaggg tgtgctcccg ggaggccagc 900
tgcagcagcc tgtgctgcgg gcggggctat gacacccaga gccgcctggt ggccttctcc 960
tgccactgcc aggtgcagtg gtgctgctac gtggagtgcc agcaatgtgt gcaggaggag 1020
cttgtgtaca cctgcaagca ctagatgggc cctgtggggt tcccgaggca gtgccaggga 1080
geettetttg agageageee tgggeagaee agggeeegee tgaeegggeg ggaagteetg 1140
acgcccttcc caggattggg cactgcggca gccccggcac agggcggggc ccacctgaag 1200
cagtgtgacc tgctgaagct gtcccggcgg cagaagcagc tctgccggag ggagcccggc 1260
ctggctgaga ccctgaggga tgctgcgcac ctcggcctgc ttgagtgcca gtttcagttc 1320
cggcatgagc gctggaactg tagcctggag ggcaggatgg gcctgctcaa gagaggcttc 1380
aaagagacag ctttcctgta cgcggtgtcc tctgccgccc tcacccacac cctggcccgg 1440
gcctgcagcg ctgggcgcat ggagcgctgc acctgtgatg actctccggg gctggagagc 1500
eggeaggeet ggeagtgggg egtgtgeggt gacaacetea agtacageae caagtttetg 1560
agcaacttcc tggggtccaa gagaggaaac aaggacctgc gggcacgggc agacgccac 1620
aatacccacg tgggcatcaa ggctgtgaag agtggcctca ggaccacgtg taagtgccat 1680
ggcgtatcag gctcctgtgc cgtgcgcacc tgctggaagc agctctcccc gttccgtgag 1740
acgggccagg tgctgaaact gcgctatgac tcggctgtca aggtgtccag tgccaccaat 1800
gaggcettgg geegeetaga getgtgggee eetgeeagge agggeageet caccaaagge 1860
ctggccccaa ggtctgggga cctggtgtac atggaggact cacccagctt ctgccggccc 1920
agcaagtact cacctggcac agcaggtagg gtgtgctccc gggaggccag ctgcagcagc 1980
ctgtgctgcg ggcggggcta tgacacccag agccgcctgg tggccttctc ctgccactgc 2040
caggtgcagt ggtgctgcta cgtggagtgc cagcaatgtg tgcaggagga gcttgtgtac 2100
acctgcaagc actag
<210> 5
<211> 1086
<212> DNA
<213> Homo sapiens
<400> 5
atgaagecee tgaggaggee cettecette atttgeecet caccaccate eccaaggete 60
acctgtctcc ctcctctcgc tctctctagc ctgaccgggc gggaagtcct gacgcccttc 120
ccaggattgg gcactgcggc agccccggca cagggcgggg cccacctgaa gcagtgtgac 180
ctgctgaagc tgtcccggcg gcagaagcag ctctgccgga gggagcccgg cctggctgag 240
accetgaggg atgetgegea ceteggeetg ettgagtgee agttteagtt eeggeatgag 300
```

```
cgctggaact gtagcctgga gggcaggatg ggcctgctca agagaggctt caaagagaca 360
gettteetgt aegeggtgte etetgeegee eteaceeaea eeetggeeeg ggeetgeage 420
gctgggcgca tggagcgctg cacctgtgat gactctccgg ggctggagag ccggcaggcc 480
tggcagtggg gcgtgtgcgg tgacaacctc aagtacagca ccaagtttct gagcaacttc 540
ctggggtcca agagaggaaa caaggacctg cgggcacggg cagacgccca caatacccac 600
gtgggcatca aggctgtgaa gagtggcctc aggaccacgt gtaagtgcca tggcgtatca 660
ggctcctgtg ccgtgcgcac ctgctggaag cagctctccc cgttccgtga gacgggccag 720
gtgctgaaac tgcgctatga ctcggctgtc aaggtgtcca gtgccaccaa tgaggccttg 780
ggccgcctag agctgtgggc ccctgccagg cagggcagcc tcaccaaagg cctggcccca 840
aggtctgggg acctggtgta catggaggac tcacccagct tctgccggcc cagcaagtac 900
tcacctggca cagcaggtag ggtgtgctcc cgggaggcca gctgcagcag cctgtgctgc 960
gggcggggct atgacaccca gagccgcctg gtggccttct cctgccactg ccaqqtqcaq 1020
tggtgctgct acgtggagtg ccagcaatgt gtgcaggagg agcttgtgta cacctgcaag 1080
cactag
                                                                   1086
<210> 6
<211> 1098
<212> DNA
<213> Homo sapiens
<400> 6
atgtggctgc ttttaacaac aacttgtttg atctgtggaa ctttaaatgc tggtggattc 60
cttgatttgg aaaatgaagt gaatcctgag gtgtggatga atactagtga aatcatcatc 120
tacaatggct accccagtga agagtatgaa gtcaccactg aagatgggta tatactcctt 180
gtcaacagaa ttccttatgg gcgaacacat gctaggagca cagcagatgc aggttatgat 240
gtatggatgg gaaacagtcg gggaaacact tggtcaagaa gacacaaaac actctcagag 300
acagatgaga aattetggge etttagtttt gatgaaatgg ecaaatatga teteceagga 360
gtaatagact tcattgtaaa taaaactggt caggagaaat tgtatttcat tggacattca 420
cttggcacta caatagggtt tgtagccttt tccaccatgc ctgaactggc acaaagaatc 480
aaaatgaatt ttgccttggg tcctacgatc tcattcaaat atcccacggg catttttacc 540
aggttttttc tacttccaaa ttccataatc aaggctgttt ttggtaccaa aggtttcttt 600
ttagaagata agaaaacgaa gatagcttct accaaaatct gcaacaataa gatactctgg 660
ttgatatgta gcgaatttat gtccttatgg gctggatcca acaagaaaaa tatgaatcag 720
agtcgaatgg atgtgtatat gtcacatgct cccactggtt catcagtaca caacattctg 780
catataaaac agctttacca ctctgatgaa ttcagagctt atgactgggg aaatgacgct 840
gataatatga aacattacaa tcagagtcat ccccctatat atgacctgac tgccatgaaa 900
gtgcctactg ctatttgggc tggtggacat gatgtcctcg taacacccca ggatgtggcc 960
aggatactcc ctcaaatcaa gagtcttcat tactttaagc tattgccaga ttggaaccac 1020
tttgattttg tctggggcct cgatgcccct caacggatgt acagtgaaat catagcttta 1080
atgaaggcat attcctaa
                                                                  1098
<210> 7
<211> 1194
<212> DNA
<213> Homo sapiens
<400> 7
atgtggcagc ttttagcagc agcatgctgg atgcttcttc ttggatctat gtatggttat 60
gacaagaaag gaaacaatgc aaaccctgaa gctaatatga atattagcca gattatttct 120
tactggggtt atccttatga agagtatgat gttacaacaa aagatggtta tatccttgga 180
atttatagga ttccacatgg aagaggatgc ccagggagga cagctccaaa gcctgctgtg 240
tatttgcagc atggcttaat tgcatctgcc agtaactgga tttgcaacct gcccaacaac 300
agtttggctt tccttctggc agatagtggt tatgacgtgt ggttggggaa cagccgagga 360
aacacttggt ccagaaaaca ccttaaattg tcaccgaaat caccggaata ctgggccttc 420
agtttggatg agatggctaa atatgacctt ccagccacaa tcaattttat catagagaaa 480
actggacaga agcgactcta ctacgtgggc cactcacaag gcaccaccat agcttttata 540
gcattttcta caaacccaga actggctaaa aagattaaga tattttttgc actggctcca 600
```

```
gttgtcacag ttaaatacac ccaaagtcct atgaaaaaac taacaaccct ttccaggcga 660
gtagttaagg tgttgtttgg tgacaaaatg ttccaccctc atacattgtt tgaccaattc 720
attgccacca aagtgtgcaa tcgaaagcta ttccgtcgta tttgcagcaa cttcctattt 780
actctgagtg gatttgatcc gcaaaactta aatatgagtc gcttggatgt ttatttgtca 840
cacaatcctg cgggaacatc tgttcagaat atgctgcact gggctcagct ttaccactct 900
gatgaattca gagcttatga ctggggaaat gacgctgata atatgaaaca ttacaatcag 960
agtcatcccc ctatatatga cctgactgcc atgaaagtgc ctactgctat ttgggctggt 1020
ggacatgatg tcctcgtaac accccaggat gtggccagga tactccctca aatcaagagt 1080
cttcattact ttaagctatt gccagattgg aaccactttg attttgtctg gggcctcgat 1140
gcccctcaac ggatgtacag tgaaatcata gctttaatga aggcatattc ctaa
<210> 8
<211> 11118
<212> DNA
<213> Homo sapiens
<400> 8
atggcgaage ggetetgege ggggagegea etgtgtgtte geggeeeeeg gggeeeegeg 60
ccgctgctgc tggtcgggct ggcgctgctg ggcgcggcgc gggcgcggga ggaggcgggc 120
ggcggcttca gcctgcaccc gccctacttc aacctggccg agggcgcccg catcgccgcc 180
teegegaeet geggagagga ggeeeeggeg egeggeteee egegeeeeae egaggaeett 240
tactgcaagc tggtaggggg ccccgtggcc ggcggcgacc ccaaccagac catccggggc 300
cagtactgtg acatctgcac ggctgccaac agcaacaagg cacaccccgc gagcaatgcc 360
atcgatggca cggagcgctg gtggcagagt ccaccgctgt cccgcggcct ggagtacaac 420
gaggtcaacg tcaccctgga cctgggccag gtcttccacg tggcctacgt cctcatcaag 480
tttgccaact caccccggcc ggacctctgg gtgctggagc ggtccatgga cttcggccgc 540
acctaccage cetggeagtt etttgeetee tecaagaggg actgtetgga geggtteggg 600
ccacagacgc tggagcgcat cacacgggac gacgcggcca tctgcaccac cgagtactca 660
cgcatcgtgc ccctggagaa cggagagatc gtggtgtccc tggtgaacgg acgtccgggc 720
gccatgaatt teteetaete geegetgeta egtgagttea ceaaggeeae eaaegteege 780
ctgcgcttcc tgcgtaccaa cacgctgctg ggccatctca tggggaaggc gctgcgggac 840
cccacggtca cccgccggta ttattacagc atcaaggata tcagcatcgg aggccgctgt 900
gtctgccacg gccacgcgga tgcctgcgat gccaaagacc ccacggaccc gttcaggctg 960
cagtgcacct gccagcacaa cacctgcggg ggcacctgcg accgctgctg ccccggcttc 1020
aatcagcagc cgtggaagcc tgcgactgcc aacagtgcca acgagtgcca gtcctgtaac 1080
tgctacggcc atgccaccga ctgttactac gaccctgagg tggaccggcg ccgcgccagc 1140
cagageetgg atggeaceta teagggtggg ggtgtetgta tegaetgeea geaceaeae 1200
accggcgtca actgtgageg ctgcctgccc ggcttctacc gctctcccaa ccaccctctc 1260
gactegeece aegtetgeeg eegetgeaac tgegagteeg aetteaegga tggeaectge 1320
gaggacetga egggtegatg etactgeegg eccaacttet etggggageg gtgtgaegtg 1380
tgtgccgagg gcttcacggg cttcccaagc tgctacccga cgccctcgtc ctccaatgac 1440
accagggage aggtgetgee ageeggeeag attgtgaatt gtgactgeag egeggeaggg 1500
acccagggca acgcctgccg gaaggaccca agggtgggac gctgtctgtg caaacccaac 1560
ttccaaggca cccattgtga gctctgcgcg ccagggttct acggccccgg ctgccagccc 1620
tgccagtgtt ccagccctgg agtggccgat gaccgctgtg accctgacac aggccagtgc 1680
aggtgccgag tgggcttcga gggggccaca tgtgatcgct gtgcccccgg ctactttcac 1740
ttccctctct gccagttgtg tggctgcagc cctgcaggaa ccttgcccga gggctgcgat 1800
gaggccggcc gctgcctatg ccagcctgag tttgctggac ctcattgtga ccggtgccgc 1860
cctggctacc atggtttccc caactgccaa gcatgcacct gcgaccctcg gggagccctg 1920
gaccagetet gtggggeggg aggtttgtge egetgeegee eeggetacae aggeaetgee 1980
tgccaggaat gcagcccgg ctttcacggc ttccccagct gtgtcccctg ccactgctct 2040
gctgaaggct ccctgcacgc agcctgtgac ccccggagtg ggcagtgcag ctgccggccc 2100
cgtgtgacgg ggctgcggtg tgacacatgt gtgcccggtg cctacaactt cccctactgc 2160
gaagetgget cttgccaccc tgccggtctg gccccagtgg atcctgccct tcctgaggca 2220
caggittccct gtatgtgccg ggctcacgtg gaggggccga gctgtgaccg ctgcaaacct 2280
gggttctggg gactgagccc cagcaacccc gagggctgta cccgctgcag ctgcgacctc 2340
aggggcacac tgggtggagt tgctgagtgc cagccgggca ccggccagtg cttctgcaag 2400
```

```
ccccacgtgt gcggccaggc ctgcgcgtcc tgcaaggatg gcttctttgg actggatcag 2460
gctgactatt ttggctgccg cagctgccgg tgtgacattg gcggtgcact gggccagagc 2520
tgtgaaccga ggacgggcgt ctgccggtgc cgccccaaca cccagggccc cacctgcagc 2580
gageetgega gggaeeacta eeteeeggae etgeaeeace tgegeetgga getggaggag 2640
gctgccacac ctgagggtca cgccgtgcgc tttggcttca acccctcga gttcgagaac 2700
ttcagctgga ggggctacgc gcagatggca cctgtccagc ccaggatcgt ggccaggctg 2760
aacctgacct cccctgacct tttctggctc gtcttccgat acgtcaaccg gggggccatg 2820
agtgtgagcg ggcgggtctc tgtgcgagag gagggcaggt cggccacctg cgccaactgc 2880
acagcacaga gtcagcccgt ggccttccca cccagcacgg agcctgcctt catcaccgtg 2940
ccccagaggg gcttcggaga gccctttgtg ctgaaccctg gcacctgggc cctgcgtgtg 3000
gaggeegaag gggtgeteet ggaetaegtg gttetgetge 🖰tagegeata etaegaggeg 3060
gcgctcctgc agctgcgggt gactgaggcc tgcacatacc gtccctctgc ccagcagtct 3120
ggcgacaact gcctcctcta cacacacctc cccctggatg gcttcccctc ggccgccggg 3180
ctggaggccc tgtgtcgcca ggacaacagc ctgccccggc cctgccccac ggagcagctc 3240
agcccgtcgc acccgccact gatcacctgc acgggcagtg atgtggacgt ccagcttcaa 3300
gtggcagtgc cacagccagg ccgctatgcc ctagtggtgg agtacgccaa tgaggatgcc 3360
cgccaggagg tgggcgtggc cgtgcacacc ccacagcggg ccccccagca ggggctgctc 3420
teeetgeace cetgeetgta cageaceetg tgeeggggea etgeeeggga tacceaggae 3480
cacctggctg tettecacet ggacteggag gecagegtga ggeteacage egaacaggea 3540
cgcttcttcc tgcacggggt cactctggtg cccattgagg agttcagccc ggagttcgtg 3600
gageceeggg teagetgeat eageageeac ggegeetttg geeceaacag tgeegeetgt 3660
ctgccctcgc gcttcccaaa gccgccccag cccatcatcc tcagggactg ccaggtgatc 3720
cegetgeege ceggeetece getgacecae gegeaggate teactecage catgteecca 3780
getggacece gaeeteggee ecceaeeget gtggaceetg atgeagagee caeeetgetg 3840
cgtgagcccc aggccaccgt ggtcttcacc acccatgtgc ccacgctggg ccgctatgcc 3900
ttcctgctgc acggctacca gccagcccac cccaccttcc ccgtggaagt cctcatcaac 3960
gccggccgcg tgtggcaggg ccacgccaac gccagcttct gtccacatgg ctacggctgc 4020
cgcaccctgg tggtgtgtga gggccaggcc ctgctggacg tgacccacag cgagctcact 4080
gtgaccgtgc gtgtgcccaa gggccggtgg ctctggctgg attatgtact cgtggtccct 4140
gagaacgtct acagctttgg ctacctccgg gaggagcccc tggataaatc ctatgacttc 4200
atcagecaet gegeageeca gggetaecae atcageecea geageteate eetgttetge 4260
cgaaacgctg ctgcttccct ctccctcttc tataacaacg gagcccgtcc atgtggctgc 4320
cacgaagtag gtgctacagg ccccacgtgt gagcccttcg ggggccagtg tccctgccat 4380
gcccatgtca ttggccgtga ctgctcccgc tgtgccaccg gatactgggg cttccccaac 4440
tgcaggccct gtgactgcgg tgcccgcctc tgtgacgagc tcacgggcca gtgcatctgc 4500
ccgccacgca ccatcccgcc cgactgcctg ctgtgccagc cccagacctt tggctgccac 4560
eccetggteg getgtgagga gtgtaactge teagggeeeg geateeagga geteaeagae 4620
cctacctgtg acacagacag cggccagtgc aagtgcagac ccaacgtgac tgggcgccgc 4680
tgtgatacct gctctccggg cttccatggc tacccccgct gccgcccctg tgactgtcac 4740
gaggcgggca ctgcgcctgg cgtgtgtgac cccctcacag ggcagtgcta ctgtaaggag 4800
aacgtgcagg gccccaaatg tgaccagtgc agccttggga ccttctcact ggatgctgcc 4860
aaccccaaag gttgcacccg ctgcttctgc tttggggcca cggagcgctg ccggagctcg 4920
tectacacce gecaggagtt egtggatatg gagggatggg tgetgetgag caetgacegg 4980
caggtggtgc cccacgagcg gcagccaggg acggagatgc tccgtgcaga cctgcggcac 5040
gtgcctgagg ctgtgcccga ggctttcccc gagctgtact ggcaggcccc accctcctac 5100
ctgggggacc gggtgtcatc ctacggtggg accetccgtt atgaactgca ctcagagacc 5160
cagcggggag atgtctttgt ccccatggag agcaggccgg atgtggtgct gcagggcaac 5220
cagatgagca tcacattcct ggagccggca taccccacgc ctggccacgt tcaccgtggg 5280
cagctgcagc tggtggaggg gaacttccgg catacggaga cgcgcaacac tgtgtcccgc 5340
gaggagetea tgatggtget ggeeageetg gageagetge agateegtge cetettetea 5400
cagateteet eggetgtett eetgegeagg gtggeaetgg aggtggeeag eeeageagge 5460
cagggggccc tggccagcaa tgtggagctg tgcctgtgcc ccgccagcta ccggggggac 5520
teatgecagg aatgtgeece eggettetat egggaegtea aaggtetett eetgggeega 5580
tgtgtccctt gtcagtgcca tggacactca gaccgctgcc tccctggctc tggcgtctgt 5640
gtggactgcc agcacaacac cgaaggggcc cactgtgagc gctgccaggc tggcttcgtg 5700
agcagcaggg acgaccccag cgcccctgt gtcagctgcc cctgccccct ctcagtgcct 5760
tccaacaact tcgccgaggg ctgtgtcctg cgaggcggcc gcacccagtg cctctgcaaa 5820
```

```
cctggttatg caggtgcctc ctgcgagcgg tgtgcgcccg gattctttgg gaacccactg 5880
gtgctgggca gctcctgcca gccatgcgac tgcagcggca acggtgaccc caacttgctc 5940
ttcagcgact gcgaccccct gacgggcgcc tgccgtggct gcctgcgcca caccactggg 6000
ccccgctgcg agatctgtgc ccccggcttc tacggcaacg ccctgctgcc cggcaactgc 6060
acceggtgeg actgtacece atgtgggaca gaggeetgeg acceecacag egggeactge 6120
ctgtgcaagg cgggcgtgac tgggcggcgc tgtgaccgct gccaggaggg acattttggt 6180
ttcgatggct gcgggggctg ccgccgtgt gcttgtggac cggccgccga gggctccgag 6240
tgccaccccc agagcggaca gtgccactgc cgaccaggga ccatgggacc ccagtgccgc 6300
gagtgtgccc ctggctactg ggggctccct gagcagggct gcaggcgctg ccagtgccct 6360
gggggccgct gtgaccctca cacgggccgc tgcaactgcc ccccggggct cagcgggag 6420
cgctgcgaca cctgcagcca gcagcatcag gtgcctgttc caggcgggcc tgtgggccac 6480
agcatccact gtgaagtgtg tgaccactgt gtggtcctgc tcctggatga cctggaacgg 6540
geoggegeee tecteeeege catteaegag caactgegtg geateaatge cageteeatg 6600
gcctgggccc gtctgcacag gctgaacgcc tccatcgctg acctgcagag ccagctccgg 6660
agccccctgg gcccccgcca tgagacggca cagcagctgg aggtgctgga gcagcagagc 6720
acaagceteg ggeaggaege aeggeggeta ggeggeeagg eaggageeee aagaeeeeee 6780
agggccccgg gaggctttca cctgtaccag gcgagccaat tgctggccgg caccgaggcc 6840
acactgggcc atgcgaagac gctgttggcg gccatccggg ctgtggaccg caccctgagc 6900
gageteatgt eccagaeggg ecacetgggg etggeeaatg ecteggetee ateaggtgag 6960
cagctgctcc ggacactggc cgaggtggag cggctgctct gggagatgcg ggcccgggac 7020
ctgggggccc cgcaggcagc agctgaggct gagttggctg cagcacagag attgctggcc 7080
cgggtgcagg agcactgag cagcctctgg gaggagaacc aggcactggc cacacaaacc 7140
cgcgaccggc tggcccagca cgaggccggc ctcatggacc tgcgagaggc tttgaaccgg 7200
gcagtggacg ccacacggga ggcccaggag ctcaacagcc gcaaccagga gcgcctggag 7260
gaagccetge aaaggaagca ggagetgtee egggacaatg ceaccetgea ggecaetetg 7320
catgcggcta gggacaccct ggccagcgtc ttcagattgc tgcacagcct ggaccaggct 7380
aaggaggage tggagegeet egeegeeage etggatgggg eteggaeeee aetgetgeag 7440
aggatgcaga cettetecce ggegggeage aagetgegte tagtggagge egeegaggee 7500
cacgcacage agetgggeea getggeacte aatetgteea geateateet ggaegteaac 7560
caggaccgcc tcacccagag ggccatcgag gcctccaacg cctacagccg catcctgcag 7620
gccgtgcagg ctgccgagga tgctgctggc caggccctgc agcaggcgga ccacacgtgg 7680
gcgacggtgg tgcggcaggg cctggtggac cgagcccagc agctcctggc caacagcact 7740
gcactagaag aggccatgct ccaggaacag cagaggctgg gccttgtgtg ggctgccctc 7800
cagggtgcca ggacccagct ccgagatgtc cgggccaaga aggaccagct ggaggcgcac 7860
atccaggcgg cgcaggccat gcttgccatg gacacagacg agacaagcaa gaagatcgca 7920
catgccaagg ctgtggctgc tgaagcccag gacaccgcca cccgtgtgca gtcccagctg 7980
caggccatgc aggagaatgt ggagcggtgg cagggccagt acgagggcct gcggggccag 8040
gacctgggcc aggcagtgct tgacgcaggc cactcagtgt ccaccctgga gaagacgctg 8100
ccccagctgc tggccaagct gagcatcctg gagaaccgtg gggtgcacaa cgccagcctg 8160
gccctgtccg ccagcattgg ccgcgtgcga gagctcattg cccaggcccg gggggctgcc 8220
agtaaggtca aggtgcccat gaagttcaac gggcgctcag gggtgcagct gcgcacccca 8280
cgggatcttg ccgaccttgc tgcctacact gccctcaagt tctacctgca gggcccagag 8340
cctgagcctg ggcagggtac cgaggatcgc tttgtgatgt acatgggcag ccgccaggcc 8400
actggggact acatgggtgt gtctctgcgt gacaagaagg tgcactgggt gtatcagctg 8460
ggtgaggcgg gccctgcagt cctaagcatc gatgaggaca ttggggagca gttcgcagct 8520
gtcagcctgg acaggactct ccagtttggc cacatgtccg tcacagtgga gagacagatg 8580
atccaggaaa ccaagggtga cacggtggcc cctggggcag aggggctgct caacctgcgg 8640
ccagacgact tcgtcttcta cgtcgggggg taccccagta ccttcacgcc ccctccctg 8700
cttcgcttcc ccggctaccg gggctgcatc gagatggaca cgctgaatga ggaggtggtc 8760
ageetetaca aettegagag gaeetteeag etggaeaegg etgtggaeag geettgtgee 8820
cgctccaagt cgaccgggga cccgtggctc acggacggct cctacctgga cggcaccggc 8880
ttcgcccgca tcagcttcga cagtcagatc agcaccacca agcgcttcga gcaggagctg 8940
eggetegtgt cetacagegg ggtgetette tteetgaage ageagageea gtteetgtge 9000
ttggccgtgc aagaaggcag cctcgtgctg ttgtatgact ttggggctgg cctgaaaaag 9060
gccgtcccac tgcagccccc accgcccctg acctcggcca gcaaggcgat ccaggtgttc 9120
ctgctggggg gcagccgcaa gcgtgtgctg gtgcgtgtgg agcgggccac ggtgtacagc 9180
gtggagcagg acaatgatct ggagctggcc gacgcctact acctgggggg cgtgccgccc 9240
```

```
gaccagetge eccegageet gegaeggete tteeceaceg gaggeteagt eegtggetge 9300
gtcaaaggca tcaaggccct gggcaagtat gtggacctca agcggctgaa cacgacaggc 9360
gtgagcgccg gctgcaccgc cgacctgctg gtggggcgcg ccatgacttt ccatggccac 9420
ggetteette geetggeget etegaaegtg geaeegetea etggeaaegt etaeteegge 9480
ttcggcttcc acagcgccca ggacagtgcc ctgctctact accgggcgtc cccggatggg 9540
ctatgccagg tgtccctgca gcagggccgt gtgagcctac agctcctgag gactgaagtg 9600
aaaactcaag cgggcttcgc cgatggtgcc ccccattacg tcgccttcta cagcaatgcc 9660
acgggagtet ggctgtatgt cgatgaccag ctccagcaga tgaagcccca ccggggacca 9720
cccccgage tecageegea geetgagggg ecceegagge teeteetggg aggeetgeet 9780
gagtctggca ccatttacaa cttcagtggc tgcatcagca acgtcttcgt gcagcggctc 9840
ctgggcccac agcgcgtatt tgatctgcag cagaacctgg gcagcgtcaa tgtgagcacg 9900
ggctgtgcac ccgccctgca agcccagacc ccgggcctgg ggcctagagg actgcaggcc 9960
accgcccgga aggcctcccg ccgcagccgt cagcccgccc ggcatcctgc ctgcatgctg 10020
cccccacacc tcaggaccac ccgagactcc taccagtttg ggggttccct gtccagtcac 10080
ctggagtttg tgggcatcct ggcccgacat aggaactggc ccagtctctc catgcacgtc 10140
ctcccgcgaa gctcccgagg cctcctcctc ttcactgccc gtctgaggcc cggcagcccc 10200
teeetggege tetteetgag caatggeeae ttegttgeae agatggaagg cetegggaet 10260
cggctccgcg cccagagccg ccagcgctcc cggcctggcc gctggcacaa ggtctccgtg 10320
cgctgggaga agaaccggat cctgctggtg acggacgggg cccgggcctg gagccaggag 10380
gggccgcacc ggcagcacca gggggcagag cacccccagc cccacaccct ctttgtgggc 10440
ggcctcccgg ccagcagcca cagctccaaa cttccggtga ccgtcgggtt cagcggctgt 10500
gtgaagagac tgaggctgca cgggaggccc ctgggggccc ccacacggat ggcaggggtc 10560
acaccctgca tcttgggccc cctggaggcg ggcctgttct tcccaggcag cgggggagtt 10620
atcactttag acctcccagg agctacactg cctgatgtgg gcctggaact ggaggtgcgg 10680
cccctggcag tcaccggact gatcttccac ttgggccagg cccggacgcc cccctacttg 10740
cagttgcagg tgaccgagaa gcaagtcctg ctgcgggcgg atgacggagc aggggagttc 10800
tecaegteag tgaecegeee eteagtgetg tgtgatggee agtggeaeeg getageggtg 10860
atgaaaagcg ggaatgtgct ccggctggag gtggacgcgc agagcaacca caccgtgggc 10920
cccttgctgg cggctgcagc tggtgcccca gccctctgt acctcggggg cctgcctgag 10980
cccatggccg tgcagccctg gcccccgcc tactgcggct gcatgaggag gctggcggtg 11040
aaccggtccc ccgtcgccat gactcgctct gtggaggtcc acggggcagt gggggccagt 11100
ggctgcccag ccgcctag
                                                                  11118
<213> Homo sapiens
```

<210> 9 <211> 11091 <212> DNA

<400> 9

atggcgaage ggctctgcgc ggggagcgca ctgtgtgttc gcggcccccg gggccccgcg 60 eegetgetge tggteggget ggegetgetg ggegeggege gggegeggga ggaggeggge 120 ggcggcttca gcctgcaccc gccctacttc aacctggccg agggcgcccg catcgccgcc 180 teegegaeet geggagagga ggeeeeggeg egeggeteee egegeeeeae egaggaeett 240 tactgcaagc tggtaggggg ccccgtggcc ggcggcgacc ccaaccagac catccggggc 300 cagtactgtg acatctgcac ggctgccaac agcaacaagg cacaccccgc gagcaatgcc 360 atcgatggca cggagcgctg gtggcagagt ccaccgctgt cccgcggcct ggagtacaac 420 gaggtcaacg tcaccctgga cctgggccag gtcttccacg tggcctacgt cctcatcaag 480 tttgccaact caccceggce ggacetetgg gtgetggage ggtecatgga etteggeege 540 acctaccage cetggeagtt etttgeetee teeaagaggg aetgtetgga geggtteggg 600 ccacagacge tggagcgcat cacacgggac gacgcggcca tctgcaccac cgagtactca 660 cgcatcgtgc ccctggagaa cggagagatc gtggtgtccc tggtgaacgg acgtccgggc 720 gccatgaatt tetectaete geegetgeta egtgagttea ecaaggeeae caaegteege 780 ctgcgcttcc tgcgtaccaa cacgctgctg ggccatctca tggggaaggc gctgcgggac 840 cccacggtca cccgccggta ttattacagc atcaaggata tcagcatcgg aggccgctgt 900 gtctgccacg gccacgcgga tgcctgcgat gccaaagacc ccacggaccc gttcaggctg 960 cagtgcacct gccagcacaa cacctgcggg ggcacctgcg accgctgctg ccccggcttc 1020 aatcagcage cgtggaagce tgcgactgce aacagtgcca acgagtgcca gtcctgtaac 1080

```
tgctacggcc atgccaccga ctgttactac gaccctgagg tggaccggcg ccgcgccagc 1140
cagageetgg atggeaceta teagggtggg ggtgtetgta tegaetgeea geaceacace 1200
accggcgtca actgtgagcg ctgcctgccc ggcttctacc gctctcccaa ccaccctctc 1260
gactegeece aegtetgeeg eegetgeaac tgegagteeg aetteaegga tggeaectge 1320
gaggacctga cgggtcgatg ctactgccgg cccaacttct ctggggagcg gtgtgacgtg 1380
tgtgccgagg gcttcacggg cttcccaagc tgctacccga cgccctcgtc ctccaatgac 1440
accagggage aggtgetgee ageeggeeag attgtgaatt gtgaetgeag egeggeaggg 1500
acccagggca acgcctgccg gaaggaccca agggtgggac gctgtctgtg caaacccaac 1560
ttccaaggca cccattgtga gctctgcgcg ccagggttct acggccccgg ctgccagccc 1620
tgccagtgtt ccagccctgg agtggccgat gaccgctgtg accctgacac aggccagtgc 1680
aggtgccgag tgggcttcga gggggccaca tgtgatcgct gtgcccccgg ctactttcac 1740
ttccctctct gccagttgtg tggctgcagc cctgcaggaa ccttgcccga gggctgcgat 1800
gaggccggcc gctgcctatg ccagcctgag tttgctggac ctcattgtga ccggtgccgc 1860
cctggctacc atggtttccc caactgccaa gcatgcacct gcgaccctcg gggagccctg 1920
gaccagetet gtggggeggg aggtttgtge egetgeegee eeggetacae aggeaetgee 1980
tgccaggaat gcagcccgg ctttcacggc ttccccagct gtgtcccctg ccactgctct 2040
gctgaagget ccctgcacge agcctgtgac ccccggagtg ggcagtgcag ctgccggccc 2100
cgtgtgacgg ggctgcggtg tgacacatgt gtgcccggtg cctacaactt cccctactgc 2160
gaagctggct cttgccaccc tgccggtctg gccccagtgg atcctgccct tcctgaggca 2220
caggttccct gtatgtgccg ggctcacgtg gaggggccga gctgtgaccg ctgcaaacct 2280
gggttctggg gactgagccc cagcaacccc gagggctgta cccgctgcag ctgcgacctc 2340
aggggcacac tgggtggagt tgctgagtgc cagccgggca ccggccagtg cttctgcaag 2400
ccccacgtgt gcggccaggc ctgcgcgtcc tgcaaggatg gcttctttgg actggatcag 2460
gctgactatt ttggctgccg cagctgccgg tgtgacattg gcggtgcact gggccagagc 2520
tgtgaaccga ggacgggcgt ctgccggtgc cgccccaaca cccagggccc cacctgcagc 2580
gagectgega gggaceacta ecteeeggae etgeaceace tgegeetgga getggaggag 2640
getgecacae etgagggtea egeegtgege tttggettea acceetega gttegagaae 2700
ttcagctgga ggggctacgc gcagatggca cctgtccagc ccaggatcgt ggccaggctg 2760
aacctgacct cccctgacct tttctggctc gtcttccgat acgtcaaccg gggggccatg 2820
agtgtgageg ggegggtete tgtgegagag gagggeaggt eggeeaeetg egeeaaetge 2880
acagcacaga gtcagcccgt ggccttccca cccagcacgg agcctgcctt catcaccgtg 2940
ccccagaggg gcttcggaga gccctttgtg ctgaaccctg gcacctgggc cctgcgtgtg 3000
gaggeegaag gggtgeteet ggaetaegtg gttetgetge etagegeata etaegaggeg 3060
gegeteetge agetgegggt gaetgaggee tgeacatace gteeetetge ceageagtet 3120
ggcgacaact gcctctcta cacacacctc cccctggatg gcttcccctc ggccgccggg 3180
ctggaggccc tgtgtcgcca ggacaacagc ctgccccggc cctgccccac ggagcagctc 3240
agecegtege accegecaet gateacetge aegggeagtg atgtggaegt ceagetteaa 3300
gtggcagtgc cacagccagg ccgctatgcc ctagtggtgg agtacgccaa tgaggatgcc 3360
cgccaggagg tgggcgtggc cgtgcacacc ccacagcggg ccccccagca ggggctgctc 3420
tecetgeace cetgeetgta cageaceetg tgeeggggea etgeeeggga taceeaggae 3480
cacctggctg tcttccacct ggactcggag gccagcgtga ggctcacagc cgaacaggca 3540
cgcttcttcc tgcacggggt cactctggtg cccattgagg agttcagccc ggagttcgtg 3600
gageceeggg teagetgeat eageageeac ggegeetttg geeceaacag tgeegeetgt 3660
ctgccctcgc gcttcccaaa gccgccccag cccatcatcc tcagggactg ccaggtgatc 3720
ccgctgccgc ccggcctccc gctgacccac gcgcaggatc tcactccagc catgtcccca 3780
gctggacccc gacctcggcc ccccaccgct gtggaccctg atgcagagcc caccctgctg 3840
cgtgagcccc aggccaccgt ggtcttcacc acccatgtgc ccacgctggg ccgctatgcc 3900
tteetgetge aeggetacea geeageeeae eecacettee eegtggaagt eeteateaae 3960
geoggeogeg tgtggcaggg ccaegecaac geoagettet gtecacatgg ctaeggetge 4020
cgcaccctgg tggtgtgtga gggccaggcc ctgctggacg tgacccacag cgagctcact 4080
gtgaccgtgc gtgtgcccaa gggccggtgg ctctggctgg attatgtact cgtggtccct 4140
gagaacgtct acagctttgg ctacctccgg gaggagcccc tggataaatc ctatgacttc 4200
atcagccact gegeageeca gggetaceae atcageecea geageteate cetgttetge 4260
cgaaacgctg ctgcttccct ctccctcttc tataacaacg gagcccgtcc atgtggctgc 4320
cacgaagtag gtgctacagg ccccacgtgt gagcccttcg ggggccagtg tccctgccat 4380
gcccatgtca ttggccgtga ctgctcccgc tgtgccaccg gatactgggg cttccccaac 4440
tgcaggccct gtgactgcgg tgcccgcctc tgtgacgagc tcacgggcca gtgcatctgc 4500
```

```
eegecacgea ceateeegee egactgeetg etgtgeeage eecagacett tggetgeeae 4560
eccetggteg getgtgagga gtgtaactge teagggeeeg geateeagga geteaeagae 4620
cctacctgtg acacagacag cggccagtgc aagtgcagac ccaacgtgac tgggcgccgc 4680
tgtgatacct gctctccggg cttccatggc tacccccgct gccgcccctg tgactgtcac 4740
gaggcgggca ctgcgcctgg cgtgtgtgac cccctcacag ggcagtgcta ctgtaaggag 4800
aacgtgcagg gccccaaatg tgaccagtgc agccttggga ccttctcact ggatgctgcc 4860
aaccccaaag gttgcacccg ctgcttctgc tttggggcca cggagcgctg ccggagctcg 4920
tectacacce gecaggagtt egtggatatg gagggatggg tgetgetgag caetgacegg 4980
caggtggtgc cccacgagcg gcagccaggg acggagatgc tccgtgcaga cctgcggcac 5040
gtgcctgagg ctgtgcccga ggctttcccc gagctgtact ggcaggcccc accctcctac 5100
ctgggggacc gggtgtcatc ctacggtggg accetecgtt atgaactgca ctcagagacc 5160
cagcggggag atgtctttgt ccccatggag agcaggccgg atgtggtgct gcagggcaac 5220
cagatgagca tcacattcct ggagccggca taccccacgc ctggccacgt tcaccgtggg 5280
cagctgcagc tggtggaggg gaacttccgg catacggaga cgcgcaacac tgtgtcccgc 5340
gaggagetea tgatggtget ggecageetg gageagetge agateegtge eetettetea 5400
cagatetect eggetgtett eetgegeagg gtggeactgg aggtggeeag eecageagge 5460
cagggggccc tggccagcaa tgtggagctg tgcctgtgcc ccgccagcta ccggggggac 5520
tcatgccagg aatgtgcccc cggcttctat cgggacgtca aaggtctctt cctgggccga 5580
tgtgtccctt gtcagtgcca tggacactca gaccgctgcc tccctggctc tggcgtctgt 5640
gtggactgcc agcacaacac cgaaggggcc cactgtgagc gctgccaggc tggcttcgtg 5700
agcagcaggg acgaccccag cgcccctgt gtcagctgcc cctgccccct ctcagtgcct 5760
tecaacaact tegeogaggg etgtgteetg egaggeggee geacecagtg eetetgeaaa 5820
cctggttatg caggtgcctc ctgcgagcgg tgtgcgcccg gattctttgg gaacccactg 5880
gtgctgggca gctcctgcca gccatgcgac tgcagcggca acggtgaccc caacttgctc 5940
ttcagcgact gcgaccccct gacgggcgcc tgccgtggct gcctgcgcca caccactggg 6000
eccegetgeg agatetgtge ecceggette taeggeaaeg ecctgetgee eggeaaetge 6060
accoggtgcg actgtacccc atgtgggaca gaggcctgcg acccccacag cgggcactgc 6120
ctgtgcaagg cgggcgtgac tgggcggcgc tgtgaccgct gccaggaggg acattttggt 6180
ttcgatggct gcgggggctg ccgccgtgt gcttgtggac cggccgccga gggctccgag 6240
tgccacccc agageggaca gtgccactgc cgaccaggga ccatgggacc ccagtgccgc 6300
gagtgtgccc ctggctactg ggggctccct gagcagggct gcaggcgctg ccagtgccct 6360
gggggccgct gtgaccctca cacgggccgc tgcaactgcc ccccggggct cagcggggag 6420
cgctgcgaca cctgcagcca gcagcatcag gtgcctgttc caggcgggcc tgtgggccac 6480
agcatccact gtgaagtgtg tgaccactgt gtggtcctgc tcctggatga cctggaacgg 6540
geeggegeee teeteeeege catteaegag caactgegtg geateaatge cageteeatg 6600
gcctgggccc gtctgcacag gctgaacgcc tccatcgctg acctgcagag ccagctccgg 6660
agccccctgg gcccccgcca tgagacggca cagcagctgg aggtgctgga gcagcagagc 6720
acaagceteg ggeaggaege aeggeggeta ggeggeeagg eageegtggg gaeeegagae 6780
caggcgagcc aattgctggc cggcaccgag gccacactgg gccatgcgaa gacgctgttg 6840
gcggccatcc gggctgtgga ccgcaccctg agcgagctca tgtcccagac gggccacctg 6900
gggctggcca atgcctcggc tccatcaggt gagcagctgc tccggacact ggccgaggtg 6960
gageggetge tetgggagat gegggeeegg gaeetggggg eeeegeagge ageagetgag 7020
getgagttgg etgeageaea gagattgetg geeegggtge aggageaget gageageete 7080
tgggaggaga accaggcact ggccacacaa acccgcgacc ggctggccca gcacgaggcc 7140
ggcctcatgg acctgcgaga ggctttgaac cgggcagtgg acgccacacg ggaggcccag 7200
gageteaaca geegeaacea ggagegeetg gaggaageee tgeaaaggaa geaggagetg 7260
tecegggaea atgecaeect geaggeeaet etgeatgegg etagggaeae eetggeeage 7320
gtcttcagat tgctgcacag cctggaccag gctaaggagg agctggagcg cctcgccgcc 7380
agcctggatg gggctcggac cccactgctg cagaggatgc agaccttctc cccggcgggc 7440
agcaagetge gtetagtgga ggeegeegag geeeaegeac agcagetggg eeagetggea 7500
ctcaatctgt ccagcatcat cctggacgtc aaccaggacc gcctcaccca gagggccatc 7560
gaggeeteea aegeetaeag eegeateetg eaggeegtge aggetgeega ggatgetget 7620
ggccaggccc tgcagcaggc ggaccacacg tgggcgacgg tggtgcggca gggcctggtg 7680
gaccgagccc agcagctcct ggccaacagc actgcactag aagaggccat gctccaggaa 7740
cagcagagge tgggcettgt gtgggetgee etceagggtg ceaggaceea geteegagat 7800
gtccgggcca agaaggacca gctggaggcg cacatccagg cggcgcaggc catgcttgcc 7860
atggacacag acgagacaag caagaagatc gcacatgcca aggctgtggc tgctgaagcc 7920
```

			ctgcaggcca			
			caggacctgg			
			ctgccccagc			
			ctggccctgt			
cgagagctca	ttgcccaggc	ccggggggct	gccagtaagg	tcaaggtgcc	catgaagttc	8220
			ccacgggatc			
			gagcctgagc			
			gccactgggg			
			ctgggtgagg			
atcgatgagg	acattgggga	gcagttcgca	gctgtcagcc	tggacaggac	tctccagttt	8520
			atgatccagg			
			cggccagacg			
gggtacccca	gtaccttcac	gccccctccc	ctgcttcgct	tccccggcta	ccggggctgc	8700
atcgagatgg	acacgctgaa	tgaggaggtg	gtcagcctct	acaacttcga	gaggaccttc	8760
cagctggaca	cggctgtgga	caggccttgt	gcccgctcca	agtcgaccgg	ggacccgtgg	8820
ctcacggacg	gctcctacct	ggacggcacc	ggcttcgccc	gcatcagctt	cgacagtcag	8880
atcagcacca	ccaagcgctt	cgagcaggag	ctgcggctcg	tgtcctacag	cggggtgctc	8940
ttcttcctga	agcagcagag	ccagttcctg	tgcttggccg	tgcaagaagg	cagcctcgtg	9000
			aaggccgtcc			
			ttcctgctgg			
			agcgtggagc			
			cccgaccagc			
			tgcgtcaaag			
			ggcgtgagcg			
			cacggcttcc			
			ggcttcggct			
			gggctatgcc			
			gtgaaaactc			
			gccacgggag			
			ccaccccccg			
			cctgagtctg			
			ctcctgggcc			
			acgggctgtg			
			gccaccgccc			
			ctgcccccac			
			cacctggagt			
			gtcctcccgc			
			ccctccctgg			
			actcggctcc			
			gtgcgctggg			
			gaggggccgc			
			ggcggcctcc			
			tgtgtgaaga			
			gtcacaccct			
			gttatcactt			
			cggcccctgg			
			ttgcagttgc			
			ttctccacgt			
			gtgatgaaaa			
			ggccccttgc			
			gagcccatgg			
			gtgaaccggt			
			agtggctgcc			11090
9-99499		~9~99999	~g cggccgcc	cagoogocca	9	7 T O D T

```
<400> 10
atgacaaaca acagcggctc caaagccgaa ctcgttgtgg gagggaaata caaactggtg 60
cggaagatcg ggtctggctc ctttggagac gtttatctgg gcatcaccac caccaacggc 120
gaggacgtag cagtgaagct ggaatctcag aaggtcaagc acccccagtt gctgtatgag 180
agcaaactct acacgattct tcaaggtggg gttggcatcc cccacatgca ctggtatggt 240
caggaaaaag acaacaatgt gctagtcatg gaccttctgg gacccagcct cgaagacctc 300
tttaatttct gttcaagaag gttcaccatg aaaactgtac ttatgttagc cgaccagatg 360
atcagcagaa ttgaatacgt gcatacaaag aattttctac accgagacat taaaccagat 420
aacttcctga tgggtactgg gcgtcactgt aataagttgt tccttattga ttttggtttg 480
gccaaaaagt acagagacaa caggaccagg caacacatac cgtacagaga agataaacac 540
ctcattggca ctgtccgata tgccagcatc aatgcacatc ttggtattga gcagagccgc 600
cgagatgaca tggaatcctt aggctacgtt ttcatgtatt ttaatagaac cagcctgccg 660
tggcaaggac taagggctat gacaaaaaaa caaaaatatg aaaagattag tgagaagaag 720
atgtccaccc ctgttgaagt tttatgtaag gggtttcctg cagaattcgc catgtacttg 780
aactactgtc gtgggctgcg ctttgaggaa gtcccagatt acatgtatct gaggcagcta 840
ttccgcattc ttttcaggac cctgaaccac caatatgact acacatttga ttggacgatg 900
ttaaagcaga aagcagcaca gcaggcagcc tcttccagtg ggcagggtca gcaggcccaa 960
acccagacag gcaagcaaac tgaaaaaaac aagaataatg tgaaagataa ctaa
<210> 11
<211> 2667
<212> DNA
<213> Homo sapiens
<400> 11
atggagtege teetgetgee ggtgetgetg etgetggeea taetgtggae geaggetgee 60
gccctcatta atctcaagta ctcggtagaa gaggagcagc gcgccgggac ggtgattgcc 120
aacgtggcca aagacgcgcg agaggcgggc ttcgcgctgg acccccggca ggcttcagcc 180
tttcgcgtgg tgtccaactc ggctccacac ctagtggaca tcaatcccag ctctggcctg 240
ctggtcacca agcagaagat tgaccgtgat ctgctgtgcc gccagagccc caagtgcatc 300
atctcgctcg aggtcatgtc cagctcaatg gaaatctgcg tgataaaggt ggagatcaag 360
gacctgaacg acaatgegee cagttteeeg geageacaga tegagetgga gateteggag 420
gcagccagcc ctggcacgcg catcccgctg gacagcgctt acgatccaga ctcaggaagc 480
tttggcgtgc agacttacga gctcacgccc aacgagctgt tcggcctgga gatcaagacg 540
cgcggcgacg gctcccgctt tgccgaactc gtggtggaaa agagcctgga ccgcgagacg 600
cagtegeact acagetteeg aateaetgeg etagaeggtg gegaeeegee gegeetggge 660
accepttages tragtateaa geteacegae tecaategaea acaacceget ettragegae 720
tecacetacg eggtgagegt gecagaaaae tegeeteeca acacaceegt cateegeete 780
aacgccagcg atccagacga gggcaccaac ggccaggtgg tctactcctt ctatggctac 840
gtcaacgacc gcacgcgcga gctctttcag atcgacccgc acagtggcct ggtcactgtc 900
actggcgctt tagactacga agaggggcac gtgtacgaac tggacgtgca ggctaaggac 960
ttggggccca attccatccc ggcacactgc aaggtcaccg tcagcgtgct ggacaccaat 1020
gacaatccgc cggtcatcaa cctgctgtca gtcaacagtg agcttgtgga ggtcagcgag 1080
agegeeecc egggetacgt gategeettg gtgegggtgt etgategega etcaggeete 1140
aatggacgtg tgcagtgccg tttgctgggc aatgtgccct ttcgactgca ggaatatgag 1200
agetteteca etattetggt ggaeggaegg etggaeegeg ageageaega eeaataeaae 1260
ctcacaattc aggcacgcga cggcggcgtg cccatgctgc agagtgccaa gtcctttacc 1320
gtgctcatca ctgacgaaaa tgacaaccac ccgcactttt ccaagcccta ctaccaggtc 1380
attgtgcagg agaacaacac gcctggcgcc tatctgctct ctgtgtctgc tcqcgacccc 1440
gacctgggtc tcaacggcag tgtctcctac cagatcgtgc cgtcgcaggt gcgggacatg 1500
cctgtcttca cctatgtctc catcaatccc aactcaggcg acatctacgc gctgcgatcc 1560
tttaaccacg agcagaccaa ggcgttcgaa ttcaaggtgc tggccaagga cggcggcctt 1620
```

coetcactge anageaacge tacggtgegg gteateatee tegacgteaa egacaacace 1680 ceggteatea eageeceace tetgattaac ggeactgeeg aggtetacat acceegeaac 1740 tetggeatag getacetggt gactgttgte aaggeagaag actacgatga gggegaaaat 1800

```
ggccgagtca cctacgacat gaccgagggc gaccgcggct tctttgaaat agaccaggtc 1860
aatggcgaag tcagaaccac ccgcaccttc ggggagagct ccaagtcctc ctatgagctt 1920
atcgtggtgg ctcacgacca cggcaagaca tctctctctg cctctgctct cgtcctaatc 1980
tacttgtccc ctgctctcga tgcccaagag tcaatgggct ctgtgaactt gtccttgatt 2040
ttcattattg ccctgggctc cattgcgggc atcctctttg taactatgat cttcgtggca 2100
atcaagtgca agcgagacaa caaagagatc cggacctaca actgcagtaa ttgtttaacc 2160
atcacttgtc tcctcggctg ttttataaaa ggacaaaaca gcaagtgtct gcattgcatc 2220
teggtttete ceattagega ggageaagae aaaaagaeag aggagaaagt gageetaagg 2280
ggaaagagaa ttgctgagta ctcctatggg catcaaaaga aatcaagcaa gaagaaaaaa 2340
atcagtaaga atgacatccg cctggtaccc cgggatgtgg aggagacaga caagatgaac 2400
gttgtcagtt gctcttccct gacctcctcc ctcaactatt ttgactacca ccagcagacg 2460
ctgcccctgg gctgccgccg ctctgagagc actttcctga atgtggagaa ccagaatacc 2520
egcaacacca gtgctaacca catctaccat cactetttca acagecaggg gccccagcag 2580
cctgacctga ttatcaacgg tgtgcctctg cctgaggtga gtgcagctaa gtggctctgt 2640
gaggttctcc caggtctcct tctttag
                                                                  2667
<210> 12
<211> 2568
<212> DNA
<213> Homo sapiens
<400> 12
atggagtege teetgetgee ggtgetgetg etgetggeea taetgtggae geaggetgee 60
geceteatta ateteaagta eteggtagaa gaggageage gegeegggae ggtgattgee 120
aacgtggcca aagacgcgcg agaggcgggc ttcgcgctgg acccccggca ggcttcagcc 180
tttcgcgtgg tgtccaactc ggctccacac ctagtggaca tcaatcccag ctctggcctg 240
ctggtcacca agcagaagat tgaccgtgat ctgctgtgcc gccagagccc caagtgcatc 300
atctcgctcg aggtcatgtc cagctcaatg gaaatctgcg tgataaaggt ggagatcaag 360
gacctgaacg acaatgcgcc cagtttcccg gcagcacaga tcgagctgga gatctcggag 420
gcagccagcc ctggcacgcg catcccgctg gacagcgctt acgatccaga ctcaggaagc 480
tttggcgtgc agacttacga gctcacgccc aacgagctgt tcggcctgga gatcaagacg 540
egeggegaeg geteeegett tgeegaaete gtggtggaaa agageetgga eegegagaeg 600
cagtcgcact acagetteeg aateaetgeg etagaeggtg gegaeeegee gegeetggge 660
accgttggcc ttagtatcaa ggtgaccgac tccaatgaca acaacccggt gtttagcgag 720
tecacetaeg eggtgagegt gecagaaaae tegeeteeca acacaceegt cateegeete 780
aacgccagcg atccagacga gggcaccaac ggccaggtgg tctactcctt ctatggctac 840
gtcaacgacc gcacgcgcga gctctttcag atcgacccgc acagtggcct ggtcactgtc 900
actggcgctt tagactacga agaggggcac gtgtacgaac tggacgtgca ggctaaggac 960
ttggggccca attccatccc ggcacactgc aaggtcaccg tcagcgtgct ggacaccaat 1020
gacaatccgc cggtcatcaa cctgctgtca gtcaacagtg agcttgtgga ggtcagcgag 1080
agegeeece egggetaegt gategeettg gtgegggtgt etgategega eteaggeete 1140
aatggacgtg tgcagtgccg tttgctgggc aatgtgccct ttcgactgca ggaatatgag 1200
agcttctcca ctattctggt ggacggacgg ctggaccgcg agcagcacga ccaatacaac 1260
ctcacaattc aggcacgcga cggcggcgtg cccatgctgc agagtgccaa gtcctttacc 1320
gtgctcatca ctgacqaaaa tgacaaccac ccgcactttt ccaaqcccta ctaccaqqtc 1380
attgtgcagg agaacaacac gcctggcgcc tatctgctct ctgtgtctgc tcgcgacccc 1440
gacctgggtc tcaacggcag tgtctcctac cagatcgtgc cgtcgcaggt gcgggacatg 1500
cctgtcttca cctatgtctc catcaatccc aactcaggcg acatctacgc gctgcgatcc 1560
tttaaccacg agcagaccaa ggcgttcgaa ttcaaggtgc tggccaagga cggcggcctt 1620
ccctcactgc aaagcaacgc tacggtgcgg gtcatcatcc tcgacgtcaa cgacaacacc 1680
ccggtcatca cagccccacc tctgattaac ggcactgccg aggtctacat accccgcaac 1740
tctggcatag gctacctggt gactgttgtc aaggcagaag actacgatga gggcgaaaat 1800
ggccgagtca cctacgacat gaccgagggc gaccgcggct tctttgaaat agaccaggtc 1860
aatggcgaag tcagaaccac ccgcaccttc ggggagagct ccaagtcctc ctatgagctt 1920
atcgtggtgg ctcacgacca cggcaagaca tctctctctg cctctgctct cgtcctaatc 1980
tacttgtccc ctgctctcga tgcccaagag tcaatgggct ctgtgaactt gtccttgatt 2040
```

ttcattattg ccctgggctc cattgcgggc atcctctttg taactatgat cttcgtggca 2100

```
atcaagtgca agcgagacaa caaagagatc cggacctaca actgcagaat tgctgagtac 2160
tcctatgggc atcaaagaa atcaagcaag aagaaaaaaa tcagtaagaa tgacatccgc 2220
ctggtacccc gggatgtgga ggagacagac aagatgaacg ttgtcagttg ctcttccctg 2280
acctectece teaactattt tgactaceae eageagaege tgeecetggg etgeegeege 2340
tetgagagea ettteetgaa tgtggagaac cagaataece geaacaecag tgetaaceae 2400
atctaccatc actetttcaa cagecagggg ceecageage etgacetgat tateaaeggt 2460
gtgcctctgc ctgagactga aaactattct tttgactcca actacgtgaa tagccgagcc 2520
catttaatca agaggtatgt tggtttgctt gcttattgct gcaactaa
<210> 13
<211> 990
<212> DNA
<213> Homo sapiens
<400> 13
atggtgacga aggcctttgt cttgttggcc atctttgcag aagcctctgc aaaatcgtgt 60
gctccaaata aagcagatgt cattcttgtg ttttgctatc ccaaaaccat catcaccaaa 120
atccccgagt gtccctatgg atgggaagtt catcagctgg ccctcggagg gctgtgttac 180
aatggggtcc acgaaggagg ttactaccaa tttgtgatcc cagatttatc acctaaaaac 240
aagtcctatt gtggaaccca gtctgagtac aagccaccta tctatcactt ctacagtcac 300
atcgtttcca atgacaccac agtgattgta aaaaaccagc ctgtcaacta ctccttctcc 360
tgcacctacc actccaccta cttggtgaac caggctgcct ttgaccagag agtggccact 420
gttcacgtga agaacgggag catgggcaca tttgagagcc aactgtctct caacttctac 480
actaatgcca agttctccat caagaaagaa gctccctttg tcctggaggc atccgaaatc 540
ggttcagatc tgtttgcagg agtggaagcc aaagggttaa gcattaggtt taaagtggtc 600
ttgaacagct gttgggccac cccctcggct gacttcatgt atcccttgca gtggcagctg 660
atcaacaagg gctgccccac ggatgaaacc gtcctcgtgc atgagaatgg gagagatcac 720
agggcaacct tccaattcaa tgctttccgg ttccagaaca tccccaaact ctccaaggtg 780
tggttacact gtgagacgtt catctgcgac agtgagaaac tctcctgccc agtgacctgc 840
gataaacgga agcgcctcct gcgagaccag accgggggag tcctggtcgt ggagctctcc 900
ctgcggagca ggggattttc cagtctctat agcttctcag atgttctcca ccacctcatc 960
atgatgttgg ggatttgtgc cgtgttatag
                                                                  990
<210> 14
<211> 699
<212> DNA
<213> Homo sapiens
<400> 14
atgctctaca caaggaaaaa cctgacctgc gcacaaacca tcaactcctc agcttttggg 60
aacttgaatg tgaccaagaa aaccaccttc attgtccatg gattcaggcc aacaggctcc 120
cctcctgttt ggatggatga cttagtaaag ggtttgctct ctgttgaaga catgaacgta 180
gttgttgttg attggaatcg aggagctaca actttaatat atacccatgc ctctagtaag 240
accagaaaag tagccatggt cttgaaggaa tttattgacc agatgttggc agaaggagct 300
tctcttgatg acatttacat gatcggagta agtctaggag cccacatatc tgggtttgtt 360
ggagagatgt acgatggatg gctggggaga attacaggcc tcgaccctgc aggcccttta 420
ttcaacggga aacctcacca agacagatta gatcccagtg atgcgcagtt tgttgatgtc 480
atccattccg acactgatgg taacgctcct ttccttgtgg cactgggcta caaggagcca 540
ttaggaaaca tagacttcta cccaaatgga ggattggatc aacctggctg ccccaaaaca 600
atattgggag gaaatgttaa ggaaatgata caggcttcct atatcttttt ccttaaaaac 660
gactctatgg acttaagttc accgaaggaa gtggaatga
                                                                  699
<210> 15
<211> 1359
<212> DNA
```

<213> Homo sapiens

```
<400> 15
atgttgagat tctacttatt catcagtttg ttgtgcttgt caagatcaga cgcagaagaa 60
acatgteett catteaceag getgagettt caeagtgeag tggttggtae gggaetaaat 120
gtgaggetga tgetetaeae aaggaaaaae etgaeetgeg cacaaaceat caacteetea 180
gettttggga acttgaatgt gaccaagaaa accaeettea ttgteeatgg atteaggeea 240
acaggeteee etectgittg gatggatgae ttagtaaagg gittgetete tgitgaagae 300
atgaacgtag ttgttgttga ttggaatcga ggagctacaa ctttaatata tacccatgcc 360
tctagtaaga ccagaaaagt agccatggtc ttgaaggaat ttattgacca gatgttggca 420
gaaggagett etettgatga catttacatg ateggagtaa gtetaggage eeacatatet 480
gggtttgttg gagagatgta cgatggatgg ctggggagaa ttacaggcct cgaccctgca 540
ggccctttat tcaacgggaa acctcaccaa gacagattag atcccagtga tgcgcagttt 600
gttgatgtca tccattccga cactgatgca ctgggctaca aggagccatt aggaaacata 660
gacttctacc caaatggagg attggatcaa cctggctgcc ccaaaacaat attgggagga 720
tttcagtatt ttaaatgtga ccaccagagg tctgtatacc tgtacctgtc ttccctgaga 780
gagagetgea ceateactge gtatecetgt gaetectace aggattatag gaatggeaag 840
tgtgtcagct gcggcacgtc acaaaaagag tcctgtcccc ttctgggcta ttatgctgat 900
aattggaaag accatctaag ggggaaagat cctccaatga cgaaggcatt ctttgacaca 960
gctgaggaga gcccattctg catgtatcat tactttgtgg atattataac atgggacaag 1020
aatgtaagaa gaggggacat taccatcaaa ttgagagaca aagctggaaa cacccacaga 1080
tccaaaatca tcagtaatga acccaccaca tttcagaaat atcaccaagt gagtctactt 1140
gcaagattta atcaagatct ggataaagtg gctgcaattt ccttgatgtt ctctacagga 1200
tototaatag goocaaggta caagotoagg attotoogaa tgaagttaag gtooottgoo 1260
catccggaga ggcctcagct gtgtcggtat gatcttgtcc tgatggaaaa cgttgaaaca 1320
gtcttccaac ctattctttg cccagagttg cagttgtaa
<210> 16
<211> 1353
<212> DNA
<213> Homo sapiens
<400> 16
atggggetee ggagecacea ceteageetg ggeettetge ttetgtttet acteeetgea 60
gagtgcctgg gagctgaggg ccggctggct ctcaagctgt tccgtgacct ctttgccaac 120
tacacaagtg ccctgagacc tgtggcagac acagaccaga ctctgaatgt gaccctggag 180
gtgacactgt cccagatcat cgacatggat gaacggaacc aggtgctgac cctgtatctg 240
tggatacggc aggagtggac agatgcctac ctacgatggg accccaatgc ctatggtggc 300
ctggatgcca tccgcatccc cagcagtctt gtgtggcggc cagacatcgt actctataac 360
aaageegaeg egeageetee aggtteegee ageaceaaeg tggteetgeg eeaegatgge 420
gccgtgcgct gggacgcgcc ggccatcacg cgcagctcgt gccgcgtgga tgtagcagcc 480
ttcccgttcg acgcccagca ctgcggcctg acgttcggct cctggactca cggcgggcac 540
caactggatg tgcggccgcg cggcgctgca gccagcctgg cggacttcgt ggagaacgtg 600
gagtggcgcg tgctgggcat gccggcgcgg cggcgcgtgc tcacctacgg ctgctgctcc 660
gagecetace eegacgteae etteaegetg etgetgegee geegegeege egeetaegtg 720
tgcaacctgc tgctgccctg cgtgctcatc tcgctgcttg cgccgctcgc cttccacctg 780
cctgccgact caggcgagaa ggtgtcgctg ggcgtcaccg tgctgctggc gctcaccgtc 840
ttccagttgc tgctggccga gagcatgcca ccggccgaga gcgtgccgct catcgggaag 900
tactacatgg ccactatgac catggtcaca ttctcaacag cactcaccat ccttatcatg 960
aacctgcatt actgtggtcc cagtgtccgc ccagtgccag cctgggctag ggccctcctg 1020
ctgggacacc tggcacgggg cctgtgcgtg cgggaaagag gggagccctg tgggcagtcc 1080
aggccacctg agttatctcc tagcccccag tcgcctgaag gaggggctgg ccccccagcg 1140
ggcccttgcc acgagccacg atgtctgtgc cgccaggaag ccctactgca ccacgtagcc 1200
accattgcca ataccttccg cagccaccqa gctgcccaqc gctgccatga ggactggaag 1260
egectggeee gtgtgatgga eegettette etggeeatet tetteteeat ggeeetggte 1320
atgagectee tggtgetggt geaggeeetg tga
                                                                  1353
```

```
<212> DNA
<213> Homo sapiens
<400> 17
atggttaagg gtgagaaagg ccccaagggc aagaagatca ccctcaaggt ggccaggaat 60
tgcatcaaaa tcacttttga tgggaaaaag cgccttgact tgagcaagat gggaattacc 120
accttcccca agtgtattct gcgccttagt gacatggacg agctggacct tagccggaat 180
cttatcagga agatccctga ctccatctcc aagttccaga acctccggtg gctggacctg 240
cacagcaact acatagacaa gctgcctgag tccattggcc agatgaccag cctgctctac 300
ctcaacgtca gcaacaaccg gctgaccagc aacgggctgc ccgtggagct gaagcaactc 360
aagaacatcc gcgctgtgaa cctaggcttg aaccacctgg acagcgtgcc caccacactg 420
ggggccctga aggagctcca cgaggtaggg ctccatgaca acctactgaa caacatcccc 480
qtqaqcatct ccaaqctccc caaqctgaaa aagctcaaca taaaqcqqaa cccctttcca 540
aagccaggtg agtcggaaat attcatagac tccatcagga ggctggagaa cttgtatgtt 600
gtggaggaga aggatctgtg tgcggcttgc ctgagaaaat gccaaaacgc ccgggacaac 660
ctgaatagaa tcaagaacat ggccacgacg acaccgagaa agaccatctt tcccaatctg 720
atctcaccca attccatggc caaggactcc tgggaagact ggaggtga
                                                                  768
<210> 18
<211> 645
<212> DNA
<213> Homo sapiens
<400> 18
atgcaggcag gaactcagtc aacgcatgag tctctgaagc ctcagagggt acaatttcag 60
tcccgaaatt ttcacaacat tttqcaatqq caqcctqqqa qqqcacttac tqqcaacaqc 120
agtgtctatt ttgtgcagta caaaatatat ggacagagac aatggaaaaa taaagaagac 180
tgttggggta ctcaagaact ctcttgtgac cttaccagtg aaacctcaga catacaggaa 240
ccttattacg ggagggtgag ggcggcctcg gctgggagct actcagaatg gagcatgacg 300
ccgcggttca ctccctggtg ggaaacaaaa atagatcctc cagtcatgaa tataacccaa 360
gtcaatggct ctttgttggt aattctccat gctccaaatt taccatatag ataccaaaag 420
gaaaaaaatg tatctataga agattactat gaactactat accgagtttt tataattaac 480
aattcactag aaaaggagca aaaggtttat gaaggggctc acagagcggt tgaaattgaa 540
getetaacae cacaetecag etactgtgta gtggetgaaa tatateagee catgttagae 600
agaagaagtc agagaagtga agagagatgt gtggaaattc catga
<210> 19
<211> 696
<212> DNA
<213> Homo sapiens
<400> 19
atgatgccta aacattgctt tctaggcttc ctcatcagtt tcttccttac tggtgtagca 60
ggaactcagt caacgcatga gtctctgaag cctcagaggg tacaatttca gtcccgaaat 120
tttcacaaca ttttgcaatg gcagcctggg agggcactta ctggcaacag cagtgtctat 180
tttgtgcagt acaaaatata tggacagaga caatggaaaa ataaagaaga ctgttggggt 240
actcaagaac tetettgtga cettaceagt gaaaceteag acatacagga acettattae 300
gggagggtga gggcggcctc ggctgggagc tactcagaat ggagcatgac gccgcggttc 360
actccctggt gggaaacaaa aatagatcct ccagtcatga atataaccca agtcaatggc 420
tctttgttgg taattctcca tgctccaaat ttaccatata gataccaaaa ggaaaaaaat 480
gtatctatag aagattacta tgaactacta taccgagttt ttataattaa caattcacta 540
gaaaaggagc aaaaggttta tgaaggggct cacagagcgg ttgaaattga agctctaaca 600
ccacactcca gctactgtgt agtggctgaa atatatcagc ccatgttaga cagaagaagt 660
cagagaagtg aagagagatg tgtggaaatt ccatga
                                                                   696
<210> 20
<211> 792
```

```
<212> DNA
<213> Homo sapiens
<400> 20
atgatgccta aacattgctt tctaggcttc ctcatcagtt tcttccttac tggtgtagca 60
ggaactcagt caacgcatga gtctctgaag cctcagaggg tacaatttca gtcccgaaat 120
tttcacaaca ttttgcaatg gcagcctggg agggcactta ctggcaacag cagtgtctat 180
tttgtgcagt acaaaatcat gttctcatgc agcatgaaaa gctctcacca gaagccaagt 240
ggatgctggc agcacatttc ttgtaacttc ccaggctgca gaacattggc taaatatgga 300
cagagacaat ggaaaaataa agaagactgt tggggtactc aagaactctc ttgtgacctt 360
accagtgaaa cctcagacat acaggaacct tattacggga gggtgagggc ggcctcggct 420
gggagctact cagaatggag catgacgccg cggttcactc cctggtggga aacaaaaata 480
gatectecag teatgaatat aacceaagte aatggetett tgttggtaat tetecatget 540
ccaaatttac catatagata ccaaaaggaa aaaaatgtat ctatagaaga ttactatgaa 600
ctactatacc gagtttttat aattaacaat tcactagaaa aggagcaaaa ggtttatgaa 660
ggggctcaca gagcggttga aattgaagct ctaacaccac actccagcta ctgtgtagtg 720
gctgaaatat atcagcccat gttagacaga agaagtcaga gaagtgaaga gagatgtgtg 780
gaaattccat ga
                                                                   792
<210> 21
<211> 780
<212> DNA
<213> Homo sapiens
<400> 21
atgtatgtat tatctccagt ggaatttata attctacaac ttttatttat tcaggccatt 60
tccagcagtt taaaaggttt cctttcagct atgagactgg ctcatagagg ctgtaatgtt 120
gatacaccag tttcaacgct cacaccagtg aagacttcag aatttgaaaa ctttaaaact 180
aaaatggtta tcacatccaa aaaagactat cctctaagta agaattttcc atattccttg 240
gaacatcttc agacttctta ctgtgggctt gtccgagttg atatgcgtat gctttgctta 300
aaaagcctta ggaaattaga cttgagtcac aaccatataa aaaagcttcc agctacaatt 360
ggagacetea tacacettea agaacetaac etgaatgaca atcacetgga gteatetagt 420
gtagccttgt gtcattctac actccagaag tcacttcgga gtttggacct cagcaagaac 480
aaaatcaagg cactccctgt gcagttttgc cagctccagg aacttaagaa tttaaaactt 540
gacgataatg aattgattca atttccttgc aagataggac aactaataaa ccttcgcttt 600
ttgtcagcag ctcgaaataa gcttccattt ttgcctagtg aatttagaaa tttatccctt 660
gaatacttgg atctttttgg aaatactttt gaacaaccaa aagtccttcc agtaataaag 720
ctgcaagcac cattaacttt attggaatct tctgcacgaa ccatattaca taataggtaa 780
<210> 22
<211> 1251
<212> DNA
<213> Homo sapiens
<400> 22
atgaagctac actgtgaggt ggaggtgatc agccggcact tgcccgcctt ggggcttagg 60
aaccggggca agggcgtccg agccgtgttg agcctctgtc agcagacttc caggagtcag 120
ccgccggtcc gagccttcct gctcatctcc accctgaagg acaagcgcgg gacccgctat 180
gagctaaggg agaacattga gcaattcttc accaaatttg tagatgaggg gaaagccact 240
gttcggttaa aggagcctcc tgtggatatc tgtctaagta aggccatttc cagcagttta 300
aaaggtttcc tttcagctat gagactggct catagaggct gtaatgttga tacaccagtt 360
tcaacgctca caccagtgaa gacttcagaa tttgaaaact ttaaaactaa aatggttatc 420
acatccaaaa aagactatcc tctaagtaag aattttccat attccttgga acatcttcag 480
acticttact gigggetigt cogagtigat atgcqtatqc tittqcttaaa aagccttagg 540
aaattagact tqaqtcacaa ccatataaaa aaqcttccaq ctacaattqq aqacctcata 600
caccttcaag aacttaacct gaatgacaat cacttgqaqt catttagtgt agccttgtgt 660
```

```
cattetacae tecagaagte actteggagt ttggacetea geaagaacaa aateaaggea 720
ctccctgtgc agttttgcca gctccaggaa cttaagaatt taaaacttga cgataatgaa 780
ttgattcaat ttccttgcaa gataggacaa ctaataaacc ttcqcttttt gtcagcaqct 840
cgaaataagc ttccattttt gcctagtgaa tttagaaatt tatcccttga atacttggat 900
ctttttggaa atacttttga acaaccaaaa gtccttccag taataaagct gcaagcacca 960
ttaactttat tggaatcttc tgcacgaacc atattacata ataggaatag gattccatat 1020
ggctctcata tcattccatt ccatctctgc caagatttgg ataccgcaaa aatttgtgtt 1080
tgtggaagat tctgtctgaa ctctttcatt caaggaacta ctaccatgaa tctgcattct 1140
gttgcccaca ctgtggtctt agtagataat ttgggtggta ctgaagcacc tattatctct 1200
tatttctgtt ctctaggctg ttatgttaat tcctctgata tgttaaagta a
<210> 23
<211> 461
<212> PRT
<213> Homo sapiens
<400> 23
Met Leu Gly Ile Trp Ile Val Ala Phe Leu Phe Phe Gly Thr Ser Arg
Gly Lys Glu Val Cys Tyr Glu Arg Leu Gly Cys Phe Lys Asp Gly Leu
                                25
Pro Trp Thr Arg Thr Phe Ser Thr Glu Leu Val Gly Leu Pro Trp Ser
                            40
Pro Glu Lys Ile Asn Thr Arg Phe Leu Leu Tyr Thr Ile His Asn Pro
                        55
Asn Ala Tyr Gln Glu Ile Ser Ala Val Asn Ser Ser Thr Ile Gln Ala
                                        75
Ser Tyr Phe Gly Thr Asp Lys Ile Thr Arg Ile Asn Ile Ala Gly Trp
                                    90
Lys Thr Asp Gly Lys Trp Gln Arg Asp Met Cys Asn Val Leu Leu Gln
            100
                                105
Leu Glu Asp Ile Asn Cys Ile Asn Leu Asp Trp Ile Asn Gly Ser Arg
                            120
                                                125
Glu Tyr Ile His Ala Val Asn Asn Leu Arg Val Val Gly Ala Glu Val
                        135
                                             140
Ala Tyr Phe Ile Asp Val Leu Met Lys Lys Phe Glu Tyr Ser Pro Ser
                    150
                                        155
Lys Val His Leu Ile Gly His Ser Leu Gly Ala His Leu Ala Gly Glu
                165
                                    170
                                                         175
Ala Gly Ser Arg Ile Pro Gly Leu Gly Arg Ile Thr Gly Leu Asp Pro
            180
                                185
                                                     190
Ala Gly Pro Phe Phe His Asn Thr Pro Lys Glu Val Arg Leu Asp Pro
                            200
Ser Asp Ala Asn Phe Val Asp Val Ile His Thr Asn Ala Ala Arg Ile
                        215
                                             220
Leu Phe Glu Leu Gly Val Gly Thr Ile Asp Ala Cys Gly His Leu Asp
                    230
                                        235
Phe Tyr Pro Asn Gly Gly Lys His Met Pro Gly Cys Glu Asp Leu Ile
                245
                                    250
Thr Pro Leu Leu Lys Phe Asn Phe Asn Ala Tyr Lys Lys Glu Met Ala
            260
                                265
                                                     270
Ser Phe Phe Asp Cys Asn His Ala Arg Ser Tyr Gln Phe Tyr Ala Glu
                            280
                                                 285
Ser Ile Leu Asn Pro Asp Ala Phe Ile Ala Tyr Pro Cys Arg Ser Tyr
                        295
                                             300
Thr Ser Phe Lys Ala Gly Thr Cys Val Gly Cys Ala Asp Leu Leu His
```

```
Arg Ile Asp Lys Ile Gly Ser His Thr Ser His Val Phe Leu Thr Leu
              325
                                330
Ser Leu Pro Phe Leu Leu Val Ser Leu Tyr Leu Gly Trp Arg His Lys
           340
                              345
Leu Ser Val Lys Leu Ser Gly Ser Glu Val Thr Gln Gly Thr Val Phe
                           360
                                               365
Leu Arg Val Gly Gly Ala Val Arg Lys Thr Gly Glu Phe Ala Ile Val
                        375
                                            380
Ser Gly Lys Leu Glu Pro Gly Met Thr Tyr Thr Lys Leu Ile Asp Ala
                   390
                                       395
Asp Val Asn Val Gly Asn Ile Thr Ser Val Gln Phe Ile Trp Lys Lys
                405
                                   410
His Leu Phe Glu Asp Ser Gln Asn Lys Leu Gly Ala Glu Met Val Ile
                               425
Asn Thr Ser Gly Lys Tyr Gly Tyr Lys Ser Thr Phe Cys Ser Gln Asp
                            440
Ile Met Gly Pro Asn Ile Leu Gln Asn Leu Lys Pro Cys
                        455
```

<211> 308

<212> PRT

<213> Homo sapiens

<400> 24

Met Pro Phe Leu Gln Leu Lys Gly Arg Ala Thr Pro Pro Ser Trp Arg 5 10 His Asp Ser Arg Ser Leu Val His Leu Leu Asp Gly Lys Glu Gly Val 25 Trp Asp Thr Thr Gly Tyr Ala Leu Gly Ser Arg Glu Ser Leu Asn Pro 40 Asp Met Gly Ile Gly Asp Pro His Gly His Ser Thr Val His Thr Arg 55 60 Glu Ala Gly Thr Ala Cys Pro Leu Gln Leu Leu Gly Ala Arg Glu Ala 70 7.5 Ser Leu Leu Ala Cys Gly Ile Cys Gln Ala Ser Gly Gln Ile Phe Ile 90 Thr Gln Thr Leu Gly Ile Lys Gly Tyr Arg Thr Val Val Ala Leu Asp 100 105 Lys Val Pro Glu Asp Val Gln Glu Tyr Ser Trp Tyr Trp Gly Ala Asn 120 125 Asp Ser Ala Gly Asn Met Ile Ile Ser His Lys Pro Pro Ser Ala Gln 135 140 Gln Pro Gly Pro Met Tyr Thr Gly Arg Glu Arg Val Asn Arg Glu Gly 150 155 Ser Leu Leu Ile Arg Pro Thr Ala Leu Asn Asp Thr Gly Asn Tyr Thr 165 170 175 Val Arg Val Val Ala Gly Asn Glu Thr Gln Arg Ala Thr Gly Trp Leu 180 185 Glu Val Leu Asp Gly Pro Asp Tyr Val Leu Leu Arg Ser Asn Pro Asp 200 205 Asp Phe Asn Gly Ile Val Thr Ala Glu Ile Gly Ser Gln Val Glu Met 215 220 Glu Cys Ile Cys Tyr Ser Phe Leu Asp Leu Lys Tyr His Trp Ile His

Asn Gly Ser Leu Leu Asn Phe Ser Asp Ala Lys Met Asn Leu Ser Ser 250 245 Leu Ala Trp Glu Gln Met Gly Arg Tyr Arg Cys Thr Val Glu Asn Pro 265 260 Val Thr Gln Leu Ile Met Tyr Met Asp Val Arg Ile Gln Ala Pro His 280 Glu Cys Ser Ser Ser Pro Pro Gly Ser Cys Phe Ala His Leu Pro Ala Ser Met Pro Cys

<210> 25

<211> 457 <212> PRT

<213> Homo sapiens

<400> 25

Met Asp Leu Ser Arg Pro Arg Trp Ser Leu Trp Arg Arg Val Phe Leu 10 Met Ala Ser Leu Leu Ala Cys Gly Ile Cys Gln Ala Ser Gly Gln Ile 25 Phe Ile Thr Gln Thr Leu Gly Ile Lys Gly Tyr Arg Thr Val Val Ala Leu Asp Lys Val Pro Glu Asp Val Gln Glu Tyr Ser Trp Tyr Trp Gly 55 Ala Asn Asp Ser Ala Gly Asn Met Ile Ile Ser His Lys Pro Pro Ser 70 75 Ala Gln Gln Pro Gly Pro Met Tyr Thr Gly Arg Glu Arg Val Asn Arg 85 90 Glu Gly Ser Leu Leu Ile Arg Pro Thr Ala Leu Asn Asp Thr Gly Asn 100 105 Tyr Thr Val Arg Val Val Ala Gly Asn Glu Thr Gln Arg Ala Thr Gly 120 125 Trp Leu Glu Val Leu Glu Leu Gly Ser Asn Leu Gly Ile Ser Val Asn 135 140 Ala Ser Ser Leu Val Glu Asn Met Asp Ser Val Ala Ala Asp Cys Leu 150 155 Thr Asn Val Thr Asn Ile Thr Trp Tyr Val Asn Asp Val Pro Thr Ser 165 170 Ser Ser Asp Arg Met Thr Ile Ser Pro Asp Gly Lys Thr Leu Val Ile 180 185 Leu Arg Val Ser Arg Tyr Asp Arg Thr Ile Gln Cys Met Ile Glu Ser 200 205 Phe Pro Glu Ile Phe Gln Arg Ser Glu Arg Ile Ser Leu Thr Val Ala 215 220 Tyr Gly Pro Asp Tyr Val Leu Leu Arg Ser Asn Pro Asp Asp Phe Asn 230 235 Gly Ile Val Thr Ala Glu Ile Gly Ser Gln Val Glu Met Glu Cys Ile 250 245 Cys Tyr Ser Phe Leu Asp Leu Lys Tyr His Trp Ile His Asn Gly Ser 260 265 Leu Leu Asn Phe Ser Asp Ala Lys Met Asn Leu Ser Ser Leu Ala Trp 280 285 Glu Gln Met Gly Arg Tyr Arg Cys Thr Val Glu Asn Pro Val Thr Gln

```
Leu Ile Met Tyr Met Asp Val Arg Ile Gln Ala Pro His Glu Cys Pro
             310
                                      315
Leu Pro Ser Gly Ile Leu Pro Val Val His Arg Asp Phe Ser Ile Ser
               325
                                   330
Gly Ser Met Val Met Phe Leu Ile Met Leu Thr Val Leu Gly Gly Val
                               345
           340
                                                    350
Tyr Ile Cys Gly Val Leu Ile His Ala Leu Ile Asn His Tyr Ser Ile
        355
                           360
Arg Cys Pro His Cys Ser Gly Thr Arg Val Gly Cys Trp Leu Gly Ala
                       375
                                            380
Gly Thr Gln Glu Pro Ala Leu Pro Pro Glu Gly Lys Gln Ser Gln Lys
                   390
                                        395
Gly Arg Asp Lys Pro Gly Thr Arg Leu Ser Gly Ile Ile Trp Gly Arg
                                    410
Gln Ile Ser Pro Gln Asp Leu Lys Leu Met Gly Ala Arg Glu Gly Leu
                                425
Glu Ser Ala Met Val Leu Asn Ser Cys Gly Val Ser Ser Ser Asn Phe
                           440
Pro Ser Leu Cys Val Tyr Lys Gly Tyr
    450
                        455
```

<211> 704

<212> PRT

<213> Homo sapiens

<400> 26

Met Leu His Asp Gly Leu Thr Ala Pro Asp Gly Cys Gly Ile Tyr Ser 10 Leu Thr Gly Arg Glu Val Leu Thr Pro Phe Pro Gly Leu Gly Thr Ala 20 25 Ala Ala Pro Ala Gln Gly Gly Ala His Leu Lys Gln Cys Asp Leu Leu 40 45 Lys Leu Ser Arg Arg Gln Lys Gln Leu Cys Arg Arg Glu Pro Gly Leu 55 60 Ala Glu Thr Leu Arg Asp Ala Ala His Leu Gly Leu Leu Glu Cys Gln 70 75 Phe Gln Phe Arg His Glu Arg Trp Asn Cys Ser Leu Glu Gly Arg Met 90 Gly Leu Leu Lys Arg Gly Phe Lys Glu Thr Ala Phe Leu Tyr Ala Val 100 105 Ser Ser Ala Ala Leu Thr His Thr Leu Ala Arg Ala Cys Ser Ala Gly 120 Arg Met Glu Arg Cys Thr Cys Asp Asp Ser Pro Gly Leu Glu Ser Arg 130 135 140 Gln Ala Trp Gln Trp Gly Val Cys Gly Asp Asn Leu Lys Tyr Ser Thr 150 155 Lys Phe Leu Ser Asn Phe Leu Gly Ser Lys Arg Gly Asn Lys Asp Leu 165 170 Arg Ala Arg Ala Asp Ala His Asn Thr His Val Gly Ile Lys Ala Val 180 190 185 Lys Ser Gly Leu Arg Thr Thr Cys Lys Cys His Gly Val Ser Gly Ser 200 205 Cys Ala Val Arg Thr Cys Trp Lys Gln Leu Ser Pro Phe Arg Glu Thr

```
Gly Gln Val Leu Lys Leu Arg Tyr Asp Ser Ala Val Lys Val Ser Ser
                  230
                                    235
225
Ala Thr Asn Glu Ala Leu Gly Arg Leu Glu Leu Trp Ala Pro Ala Arg
                245
                                    250
Gln Gly Ser Leu Thr Lys Gly Leu Ala Pro Arg Ser Gly Asp Leu Val
           260
                               265
                                                    270
Tyr Met Glu Asp Ser Pro Ser Phe Cys Arg Pro Ser Lys Tyr Ser Pro
                            280
Gly Thr Ala Gly Arg Val Cys Ser Arg Glu Ala Ser Cys Ser Ser Leu
                        295
Cys Cys Gly Arg Gly Tyr Asp Thr Gln Ser Arg Leu Val Ala Phe Ser
                    310
                                        315
Cys His Cys Gln Val Gln Trp Cys Cys Tyr Val Glu Cys Gln Gln Cys
                                    330
Val Gln Glu Glu Leu Val Tyr Thr Cys Lys His Ala Met Gly Pro Val
                                345
Gly Phe Pro Arg Gln Cys Gln Gly Ala Phe Phe Glu Ser Ser Pro Gly
                            360
                                                365
Gln Thr Arg Ala Arg Leu Thr Gly Arg Glu Val Leu Thr Pro Phe Pro
                        375
                                            380
Gly Leu Gly Thr Ala Ala Pro Ala Gln Gly Gly Ala His Leu Lys
                    390
                                        395
Gln Cys Asp Leu Leu Lys Leu Ser Arg Arg Gln Lys Gln Leu Cys Arg
                                    410
Arg Glu Pro Gly Leu Ala Glu Thr Leu Arg Asp Ala Ala His Leu Gly
                                425
Leu Leu Glu Cys Gln Phe Gln Phe Arg His Glu Arg Trp Asn Cys Ser
                            440
Leu Glu Gly Arg Met Gly Leu Leu Lys Arg Gly Phe Lys Glu Thr Ala
                       455
                                            460
Phe Leu Tyr Ala Val Ser Ser Ala Ala Leu Thr His Thr Leu Ala Arg
                   470
                                       475
Ala Cys Ser Ala Gly Arg Met Glu Arg Cys Thr Cys Asp Asp Ser Pro
               485
                                    490
Gly Leu Glu Ser Arg Gln Ala Trp Gln Trp Gly Val Cys Gly Asp Asn
                                505
           500
Leu Lys Tyr Ser Thr Lys Phe Leu Ser Asn Phe Leu Gly Ser Lys Arg
                            520
Gly Asn Lys Asp Leu Arg Ala Arg Ala Asp Ala His Asn Thr His Val
                        535
                                            540
Gly Ile Lys Ala Val Lys Ser Gly Leu Arg Thr Thr Cys Lys Cys His
                    550
                                        555
Gly Val Ser Gly Ser Cys Ala Val Arg Thr Cys Trp Lys Gln Leu Ser
                565
                                    570
Pro Phe Arg Glu Thr Gly Gln Val Leu Lys Leu Arg Tyr Asp Ser Ala
           580
                                585
                                                    590
Val Lys Val Ser Ser Ala Thr Asn Glu Ala Leu Gly Arg Leu Glu Leu
                           600
Trp Ala Pro Ala Arg Gln Gly Ser Leu Thr Lys Gly Leu Ala Pro Arg
                       615
                                            620
Ser Gly Asp Leu Val Tyr Met Glu Asp Ser Pro Ser Phe Cys Arg Pro
                   630
                                        635
Ser Lys Tyr Ser Pro Gly Thr Ala Gly Arg Val Cys Ser Arg Glu Ala
                                   650
Ser Cys Ser Ser Leu Cys Cys Gly Arg Gly Tyr Asp Thr Gln Ser Arg
```

Leu Val Ala Phe Ser Cys His Cys Gln Val Gln Trp Cys Cys Tyr Val 675 680 685 Glu Cys Gln Gln Cys Val Gln Glu Glu Leu Val Tyr Thr Cys Lys His 690 695 700

<210> 27 <211> 361 <212> PRT <213> Homo sapiens

<400> 27 Met Lys Pro Leu Arg Arg Pro Leu Pro Phe Ile Cys Pro Ser Pro Pro Ser Pro Arg Leu Thr Cys Leu Pro Pro Leu Ala Leu Ser Ser Leu Thr 25 Gly Arg Glu Val Leu Thr Pro Phe Pro Gly Leu Gly Thr Ala Ala Ala 40 Pro Ala Gln Gly Gly Ala His Leu Lys Gln Cys Asp Leu Leu Lys Leu 55 Ser Arg Arg Gln Lys Gln Leu Cys Arg Arg Glu Pro Gly Leu Ala Glu 70 7.5 Thr Leu Arg Asp Ala Ala His Leu Gly Leu Leu Glu Cys Gln Phe Gln 90 85 Phe Arg His Glu Arg Trp Asn Cys Ser Leu Glu Gly Arg Met Gly Leu 105 Leu Lys Arg Gly Phe Lys Glu Thr Ala Phe Leu Tyr Ala Val Ser Ser 120 Ala Ala Leu Thr His Thr Leu Ala Arg Ala Cys Ser Ala Gly Arg Met 135 Glu Arg Cys Thr Cys Asp Asp Ser Pro Gly Leu Glu Ser Arg Gln Ala 150 155 Trp Gln Trp Gly Val Cys Gly Asp Asn Leu Lys Tyr Ser Thr Lys Phe 165 170 175 Leu Ser Asn Phe Leu Gly Ser Lys Arg Gly Asn Lys Asp Leu Arg Ala 180 185 190 Arg Ala Asp Ala His Asn Thr His Val Gly Ile Lys Ala Val Lys Ser 200 Gly Leu Arg Thr Thr Cys Lys Cys His Gly Val Ser Gly Ser Cys Ala 215 220 Val Arg Thr Cys Trp Lys Gln Leu Ser Pro Phe Arg Glu Thr Gly Gln 230 235 Val Leu Lys Leu Arg Tyr Asp Ser Ala Val Lys Val Ser Ser Ala Thr 250 245 Asn Glu Ala Leu Gly Arg Leu Glu Leu Trp Ala Pro Ala Arg Gln Gly 260 265 270 Ser Leu Thr Lys Gly Leu Ala Pro Arg Ser Gly Asp Leu Val Tyr Met 280 285 Glu Asp Ser Pro Ser Phe Cys Arg Pro Ser Lys Tyr Ser Pro Gly Thr 295 300 Ala Gly Arg Val Cys Ser Arg Glu Ala Ser Cys Ser Ser Leu Cys Cys 310 315 320 Gly Arg Gly Tyr Asp Thr Gln Ser Arg Leu Val Ala Phe Ser Cys His 330 Cys Gln Val Gln Trp Cys Cys Tyr Val Glu Cys Gln Gln Cys Val Gln

```
<211> 365
<212> PRT
<213> Homo sapiens
<400> 28
Met Trp Leu Leu Thr Thr Thr Cys Leu Ile Cys Gly Thr Leu Asn
                                10
Ala Gly Gly Phe Leu Asp Leu Glu Asn Glu Val Asn Pro Glu Val Trp
Met Asn Thr Ser Glu Ile Ile Ile Tyr Asn Gly Tyr Pro Ser Glu Glu
Tyr Glu Val Thr Thr Glu Asp Gly Tyr Ile Leu Leu Val Asn Arg Ile
                        55
Pro Tyr Gly Arg Thr His Ala Arg Ser Thr Ala Asp Ala Gly Tyr Asp
                    70
                                        75
Val Trp Met Gly Asn Ser Arg Gly Asn Thr Trp Ser Arg Arg His Lys
                                    90
Thr Leu Ser Glu Thr Asp Glu Lys Phe Trp Ala Phe Ser Phe Asp Glu
                                105
Met Ala Lys Tyr Asp Leu Pro Gly Val Ile Asp Phe Ile Val Asn Lys
                            120
Thr Gly Gln Glu Lys Leu Tyr Phe Ile Gly His Ser Leu Gly Thr Thr
                       135
Ile Gly Phe Val Ala Phe Ser Thr Met Pro Glu Leu Ala Gln Arg Ile
                   150
                                       155
Lys Met Asn Phe Ala Leu Gly Pro Thr Ile Ser Phe Lys Tyr Pro Thr
               165
                                    170
Gly Ile Phe Thr Arg Phe Phe Leu Leu Pro Asn Ser Ile Ile Lys Ala
           180
                               185
Val Phe Gly Thr Lys Gly Phe Phe Leu Glu Asp Lys Lys Thr Lys Ile
                           200
Ala Ser Thr Lys Ile Cys Asn Asn Lys Ile Leu Trp Leu Ile Cys Ser
                       215
                                            220
Glu Phe Met Ser Leu Trp Ala Gly Ser Asn Lys Lys Asn Met Asn Gln
                    230
                                        235
Ser Arg Met Asp Val Tyr Met Ser His Ala Pro Thr Gly Ser Ser Val
                                    250
His Asn Ile Leu His Ile Lys Gln Leu Tyr His Ser Asp Glu Phe Arg
                                265
Ala Tyr Asp Trp Gly Asn Asp Ala Asp Asn Met Lys His Tyr Asn Gln
                            280
                                                285
Ser His Pro Pro Ile Tyr Asp Leu Thr Ala Met Lys Val Pro Thr Ala
                       295
                                            300
Ile Trp Ala Gly Gly His Asp Val Leu Val Thr Pro Gln Asp Val Ala
                    310
                                        315
Arg Ile Leu Pro Gln Ile Lys Ser Leu His Tyr Phe Lys Leu Leu Pro
                325
                                    330
Asp Trp Asn His Phe Asp Phe Val Trp Gly Leu Asp Ala Pro Gln Arg
                               345
Met Tyr Ser Glu Ile Ile Ala Leu Met Lys Ala Tyr Ser
                            360
```

<210> 29 <211> 397 <212> PRT <213> Homo sapiens <400> 29 Met Trp Gln Leu Leu Ala Ala Cys Trp Met Leu Leu Gly Ser Met Tyr Gly Tyr Asp Lys Lys Gly Asn Asn Ala Asn Pro Glu Ala Asn Met Asn Ile Ser Gln Ile Ile Ser Tyr Trp Gly Tyr Pro Tyr Glu Glu Tyr Asp Val Thr Thr Lys Asp Gly Tyr Ile Leu Gly Ile Tyr Arg Ile Pro His Gly Arg Gly Cys Pro Gly Arg Thr Ala Pro Lys Pro Ala Val Tyr Leu Gln His Gly Leu Ile Ala Ser Ala Ser Asn Trp Ile Cys Asn Leu Pro Asn Asn Ser Leu Ala Phe Leu Leu Ala Asp Ser Gly Tyr Asp Val Trp Leu Gly Asn Ser Arg Gly Asn Thr Trp Ser Arg Lys His Leu Lys Leu Ser Pro Lys Ser Pro Glu Tyr Trp Ala Phe Ser Leu Asp Glu Met Ala Lys Tyr Asp Leu Pro Ala Thr Ile Asn Phe Ile Ile Glu Lys Thr Gly Gln Lys Arg Leu Tyr Tyr Val Gly His Ser Gln Gly Thr Thr Ile Ala Phe Ile Ala Phe Ser Thr Asn Pro Glu Leu Ala Lys Lys Ile Lys Ile Phe Phe Ala Leu Ala Pro Val Val Thr Val Lys Tyr Thr Gln Ser Pro Met Lys Lys Leu Thr Thr Leu Ser Arg Arg Val Val Lys Val Leu Phe Gly Asp Lys Met Phe His Pro His Thr Leu Phe Asp Gln Phe Ile Ala Thr Lys Val Cys Asn Arg Lys Leu Phe Arg Arg Ile Cys Ser Asn Phe Leu Phe Thr Leu Ser Gly Phe Asp Pro Gln Asn Leu Asn Met Ser Arg Leu Asp Val Tyr Leu Ser His Asn Pro Ala Gly Thr Ser Val Gln Asn Met Leu His Trp Ala Gln Leu Tyr His Ser Asp Glu Phe Arg Ala Tyr Asp Trp Gly Asn Asp Ala Asp Asn Met Lys His Tyr Asn Gln Ser His Pro Pro Ile Tyr Asp Leu Thr Ala Met Lys Val Pro Thr Ala Ile Trp Ala Gly Gly His Asp Val Leu Val Thr Pro Gln Asp Val Ala Arg Ile Leu Pro Gln Ile Lys Ser Leu His Tyr Phe Lys Leu Leu Pro Asp Trp Asn His Phe Asp Phe Val Trp Gly Leu Asp Ala Pro Gln Arg

Met Tyr Ser Glu Ile Ile Ala Leu Met Lys Ala Tyr Ser

```
<211> 3705
<212> PRT
<213> Homo sapiens
<400> 30
Met Ala Lys Arg Leu Cys Ala Gly Ser Ala Leu Cys Val Arg Gly Pro
                                    10
Arg Gly Pro Ala Pro Leu Leu Leu Val Gly Leu Ala Leu Leu Gly Ala
                                25
Ala Arg Ala Arg Glu Glu Ala Gly Gly Phe Ser Leu His Pro Pro
Tyr Phe Asn Leu Ala Glu Gly Ala Arg Ile Ala Ala Ser Ala Thr Cys
Gly Glu Glu Ala Pro Ala Arg Gly Ser Pro Arg Pro Thr Glu Asp Leu
                    70
                                        75
Tyr Cys Lys Leu Val Gly Gly Pro Val Ala Gly Gly Asp Pro Asn Gln
                                    90
Thr Ile Arg Gly Gln Tyr Cys Asp Ile Cys Thr Ala Ala Asn Ser Asn
            100
                                105
Lys Ala His Pro Ala Ser Asn Ala Ile Asp Gly Thr Glu Arg Trp Trp
                            120
                                                125
Gln Ser Pro Pro Leu Ser Arg Gly Leu Glu Tyr Asn Glu Val Asn Val
                        135
                                            140
Thr Leu Asp Leu Gly Gln Val Phe His Val Ala Tyr Val Leu Ile Lys
                   150
                                        155
Phe Ala Asn Ser Pro Arg Pro Asp Leu Trp Val Leu Glu Arg Ser Met
                                    170
Asp Phe Gly Arg Thr Tyr Gln Pro Trp Gln Phe Phe Ala Ser Ser Lys
           180
                                185
Arg Asp Cys Leu Glu Arg Phe Gly Pro Gln Thr Leu Glu Arg Ile Thr
                            200
                                                205
Arg Asp Ala Ala Ile Cys Thr Thr Glu Tyr Ser Arg Ile Val Pro
                        215
                                            220
Leu Glu Asn Gly Glu Ile Val Val Ser Leu Val Asn Gly Arg Pro Gly
                   230
                                        235
Ala Met Asn Phe Ser Tyr Ser Pro Leu Leu Arg Glu Phe Thr Lys Ala
               245
                                    250
Thr Asn Val Arg Leu Arg Phe Leu Arg Thr Asn Thr Leu Leu Gly His
                                265
Leu Met Gly Lys Ala Leu Arg Asp Pro Thr Val Thr Arg Arg Tyr Tyr
                            280
Tyr Ser Ile Lys Asp Ile Ser Ile Gly Gly Arg Cys Val Cys His Gly
    290
                        295
                                            300
His Ala Asp Ala Cys Asp Ala Lys Asp Pro Thr Asp Pro Phe Arg Leu
                    310
                                        315
Gln Cys Thr Cys Gln His Asn Thr Cys Gly Gly Thr Cys Asp Arg Cys
                325
                                    330
Cys Pro Gly Phe Asn Gln Gln Pro Trp Lys Pro Ala Thr Ala Asn Ser
           340
                                345
Ala Asn Glu Cys Gln Ser Cys Asn Cys Tyr Gly His Ala Thr Asp Cys
                           360
                                                365
Tyr Tyr Asp Pro Glu Val Asp Arg Arg Arg Ala Ser Gln Ser Leu Asp
```

```
Gly Thr Tyr Gln Gly Gly Gly Val Cys Ile Asp Cys Gln His His Thr
                  390
                                       395
Thr Gly Val Asn Cys Glu Arg Cys Leu Pro Gly Phe Tyr Arg Ser Pro
                405
                                    410
Asn His Pro Leu Asp Ser Pro His Val Cys Arg Arg Cys Asn Cys Glu
            420
                                425
Ser Asp Phe Thr Asp Gly Thr Cys Glu Asp Leu Thr Gly Arg Cys Tyr
                           440
Cys Arg Pro Asn Phe Ser Gly Glu Arg Cys Asp Val Cys Ala Glu Gly
                       455
                                            460
Phe Thr Gly Phe Pro Ser Cys Tyr Pro Thr Pro Ser Ser Ser Asn Asp
                    470
                                        475
Thr Arg Glu Gln Val Leu Pro Ala Gly Gln Ile Val Asn Cys Asp Cys
                                    490
                485
Ser Ala Ala Gly Thr Gln Gly Asn Ala Cys Arg Lys Asp Pro Arg Val
                                505
Gly Arg Cys Leu Cys Lys Pro Asn Phe Gln Gly Thr His Cys Glu Leu
                            520
Cys Ala Pro Gly Phe Tyr Gly Pro Gly Cys Gln Pro Cys Gln Cys Ser
                        535
Ser Pro Gly Val Ala Asp Asp Arg Cys Asp Pro Asp Thr Gly Gln Cys
                    550
                                        555
Arg Cys Arg Val Gly Phe Glu Gly Ala Thr Cys Asp Arg Cys Ala Pro
                                    570
                565
Gly Tyr Phe His Phe Pro Leu Cys Gln Leu Cys Gly Cys Ser Pro Ala
                                585
Gly Thr Leu Pro Glu Gly Cys Asp Glu Ala Gly Arg Cys Leu Cys Gln
Pro Glu Phe Ala Gly Pro His Cys Asp Arg Cys Arg Pro Gly Tyr His
                        615
Gly Phe Pro Asn Cys Gln Ala Cys Thr Cys Asp Pro Arg Gly Ala Leu
                   630
                                        635
Asp Gln Leu Cys Gly Ala Gly Gly Leu Cys Arg Cys Arg Pro Gly Tyr
                645
                                    650
Thr Gly Thr Ala Cys Gln Glu Cys Ser Pro Gly Phe His Gly Phe Pro
            660
                                665
Ser Cys Val Pro Cys His Cys Ser Ala Glu Gly Ser Leu His Ala Ala
                            680
Cys Asp Pro Arg Ser Gly Gln Cys Ser Cys Arg Pro Arg Val Thr Gly
                        695
                                            700
Leu Arg Cys Asp Thr Cys Val Pro Gly Ala Tyr Asn Phe Pro Tyr Cys
                    710
                                        715
Glu Ala Gly Ser Cys His Pro Ala Gly Leu Ala Pro Val Asp Pro Ala
                725
                                    730
Leu Pro Glu Ala Gln Val Pro Cys Met Cys Arg Ala His Val Glu Gly
            740
                                745
Pro Ser Cys Asp Arg Cys Lys Pro Gly Phe Trp Gly Leu Ser Pro Ser
                            760
                                                765
Asn Pro Glu Gly Cys Thr Arg Cys Ser Cys Asp Leu Arg Gly Thr Leu
                        775
                                            780
Gly Gly Val Ala Glu Cys Gln Pro Gly Thr Gly Gln Cys Phe Cys Lys
                    790
                                        795
Pro His Val Cys Gly Gln Ala Cys Ala Ser Cys Lys Asp Gly Phe Phe
                                   810
Gly Leu Asp Gln Ala Asp Tyr Phe Gly Cys Arg Ser Cys Arg Cys Asp
```

```
Ile Gly Gly Ala Leu Gly Gln Ser Cys Glu Pro Arg Thr Gly Val Cys
       835
                        840
Arg Cys Arg Pro Asn Thr Gln Gly Pro Thr Cys Ser Glu Pro Ala Arg
   850
                     855
                                        860
Asp His Tyr Leu Pro Asp Leu His His Leu Arg Leu Glu Leu Glu Glu
               870
                                   875
Ala Ala Thr Pro Glu Gly His Ala Val Arg Phe Gly Phe Asn Pro Leu
             885
                                 890
Glu Phe Glu Asn Phe Ser Trp Arg Gly Tyr Ala Gln Met Ala Pro Val
          900
                             905
Gln Pro Arg Ile Val Ala Arg Leu Asn Leu Thr Ser Pro Asp Leu Phe
       915
                         920
                                            925
Trp Leu Val Phe Arg Tyr Val Asn Arg Gly Ala Met Ser Val Ser Gly
                      935
Arg Val Ser Val Arg Glu Glu Gly Arg Ser Ala Thr Cys Ala Asn Cys
                  950
                                     955
Thr Ala Gln Ser Gln Pro Val Ala Phe Pro Pro Ser Thr Glu Pro Ala
              965
                                 970
Phe Ile Thr Val Pro Gln Arg Gly Phe Gly Glu Pro Phe Val Leu Asn
                             985
           980
                                                990
Pro Gly Thr Trp Ala Leu Arg Val Glu Ala Glu Gly Val Leu Leu Asp
                         1000
                                            1005
Tyr Val Val Leu Leu Pro Ser Ala Tyr Tyr Glu Ala Ala Leu Leu Gln
                      1015
                                        1020
Leu Arg Val Thr Glu Ala Cys Thr Tyr Arg Pro Ser Ala Gln Gln Ser
                 1030
                                    1035
Gly Asp Asn Cys Leu Leu Tyr Thr His Leu Pro Leu Asp Gly Phe Pro
                                1050
Ser Ala Ala Gly Leu Glu Ala Leu Cys Arg Gln Asp Asn Ser Leu Pro
          1060
                            1065
Arg Pro Cys Pro Thr Glu Gln Leu Ser Pro Ser His Pro Pro Leu Ile
                         1080
      1075
                                           1085
Thr Cys Thr Gly Ser Asp Val Asp Val Gln Leu Gln Val Ala Val Pro
                     1095
                                        1100
Gln Pro Gly Arg Tyr Ala Leu Val Val Glu Tyr Ala Asn Glu Asp Ala
                 1110
                                    1115
Arg Gln Glu Val Gly Val Ala Val His Thr Pro Gln Arg Ala Pro Gln
                                1130
              1125
Gln Gly Leu Leu Ser Leu His Pro Cys Leu Tyr Ser Thr Leu Cys Arg
          1140
                             1145
                                                1150
Gly Thr Ala Arg Asp Thr Gln Asp His Leu Ala Val Phe His Leu Asp
                         1160
                                            1165
Ser Glu Ala Ser Val Arg Leu Thr Ala Glu Gln Ala Arg Phe Phe Leu
   1170
                     1175
                                        1180
His Gly Val Thr Leu Val Pro Ile Glu Glu Phe Ser Pro Glu Phe Val
                 1190
                                    1195
                                                       1200
Glu Pro Arg Val Ser Cys Ile Ser Ser His Gly Ala Phe Gly Pro Asn
              1205
                                1210
                                                    1215
Ser Ala Ala Cys Leu Pro Ser Arg Phe Pro Lys Pro Pro Gln Pro Ile
         1220
                             1225
                                                1230
Ile Leu Arg Asp Cys Gln Val Ile Pro Leu Pro Pro Gly Leu Pro Leu
                         1240
                                            1245
      1235
Thr His Ala Gln Asp Leu Thr Pro Ala Met Ser Pro Ala Gly Pro Arg
                    1255
                                        1260
Pro Arg Pro Pro Thr Ala Val Asp Pro Asp Ala Glu Pro Thr Leu Leu
                                     1275
```

Arg Glu Pro Gln Ala Thr Val Val Phe Thr Thr His Val Pro Thr Leu 1285 1290 Gly Arg Tyr Ala Phe Leu Leu His Gly Tyr Gln Pro Ala His Pro Thr 1300 1305 Phe Pro Val Glu Val Leu Ile Asn Ala Gly Arg Val Trp Gln Gly His 1315 1320 1325 Ala Asn Ala Ser Phe Cys Pro His Gly Tyr Gly Cys Arg Thr Leu Val 1330 1335 1340 Val Cys Glu Gly Gln Ala Leu Leu Asp Val Thr His Ser Glu Leu Thr 1350 1355 Val Thr Val Arg Val Pro Lys Gly Arg Trp Leu Trp Leu Asp Tyr Val 1365 1370 1375 Leu Val Val Pro Glu Asn Val Tyr Ser Phe Gly Tyr Leu Arg Glu Glu 1380 1385 1390 Pro Leu Asp Lys Ser Tyr Asp Phe Ile Ser His Cys Ala Ala Gln Gly 1400 1405 Tyr His Ile Ser Pro Ser Ser Ser Leu Phe Cys Arg Asn Ala Ala 1415 1420 Ala Ser Leu Ser Leu Phe Tyr Asn Asn Gly Ala Arg Pro Cys Gly Cys 1430 1435 His Glu Val Gly Ala Thr Gly Pro Thr Cys Glu Pro Phe Gly Gly Gln 1450 1445 Cys Pro Cys His Ala His Val Ile Gly Arg Asp Cys Ser Arg Cys Ala 1460 1465 1470 Thr Gly Tyr Trp Gly Phe Pro Asn Cys Arg Pro Cys Asp Cys Gly Ala 1475 1480 1485 Arg Leu Cys Asp Glu Leu Thr Gly Gln Cys Ile Cys Pro Pro Arg Thr 1495 1500 Ile Pro Pro Asp Cys Leu Leu Cys Gln Pro Gln Thr Phe Gly Cys His 1510 1515 Pro Leu Val Gly Cys Glu Glu Cys Asn Cys Ser Gly Pro Gly Ile Gln 1525 1530 Glu Leu Thr Asp Pro Thr Cys Asp Thr Asp Ser Gly Gln Cys Lys Cys 1540 1545 1550 Arg Pro Asn Val Thr Gly Arg Arg Cys Asp Thr Cys Ser Pro Gly Phe 1555 1560 1565 His Gly Tyr Pro Arg Cys Arg Pro Cys Asp Cys His Glu Ala Gly Thr 1575 1580 Ala Pro Gly Val Cys Asp Pro Leu Thr Gly Gln Cys Tyr Cys Lys Glu 1590 1595 Asn Val Gln Gly Pro Lys Cys Asp Gln Cys Ser Leu Gly Thr Phe Ser 1605 1610 1615 Leu Asp Ala Ala Asn Pro Lys Gly Cys Thr Arg Cys Phe Cys Phe Gly 1620 1625 Ala Thr Glu Arg Cys Arg Ser Ser Ser Tyr Thr Arg Gln Glu Phe Val 1635 1640 1645 Asp Met Glu Gly Trp Val Leu Leu Ser Thr Asp Arg Gln Val Val Pro 1655 1660 His Glu Arg Gln Pro Gly Thr Glu Met Leu Arg Ala Asp Leu Arg His 1670 1675 Val Pro Glu Ala Val Pro Glu Ala Phe Pro Glu Leu Tyr Trp Gln Ala 1685 1690 Pro Pro Ser Tyr Leu Gly Asp Arg Val Ser Ser Tyr Gly Gly Thr Leu 1700 1705 1710 Arg Tyr Glu Leu His Ser Glu Thr Gln Arg Gly Asp Val Phe Val Pro 1720

```
Met Glu Ser Arg Pro Asp Val Val Leu Gln Gly Asn Gln Met Ser Ile
   1730 1735
                        1740
Thr Phe Leu Glu Pro Ala Tyr Pro Thr Pro Gly His Val His Arg Gly
         1750
                       1755
Gln Leu Gln Leu Val Glu Gly Asn Phe Arg His Thr Glu Thr Arg Asn
           1765 1770
Thr Val Ser Arg Glu Glu Leu Met Met Val Leu Ala Ser Leu Glu Gln
        1780
               1785 1790
Leu Gln Ile Arg Ala Leu Phe Ser Gln Ile Ser Ser Ala Val Phe Leu
            1800
                             1805
Arg Arg Val Ala Leu Glu Val Ala Ser Pro Ala Gly Gln Gly Ala Leu
         1815
                                   1820
Ala Ser Asn Val Glu Leu Cys Leu Cys Pro Ala Ser Tyr Arg Gly Asp
                1830
                                1835
Ser Cys Gln Glu Cys Ala Pro Gly Phe Tyr Arg Asp Val Lys Gly Leu
            1845
                             1850
Phe Leu Gly Arg Cys Val Pro Cys Gln Cys His Gly His Ser Asp Arg
         1860
                         1865
                                          1870
Cys Leu Pro Gly Ser Gly Val Cys Val Asp Cys Gln His Asn Thr Glu
                      1880
                                       1885
Gly Ala His Cys Glu Arg Cys Gln Ala Gly Phe Val Ser Ser Arg Asp
                  1895 1900
Asp Pro Ser Ala Pro Cys Val Ser Cys Pro Cys Pro Leu Ser Val Pro
               1910
                               1915
Ser Asn Asn Phe Ala Glu Gly Cys Val Leu Arg Gly Gly Arg Thr Gln
                            1930
Cys Leu Cys Lys Pro Gly Tyr Ala Gly Ala Ser Cys Glu Arg Cys Ala
                         1945
Pro Gly Phe Phe Gly Asn Pro Leu Val Leu Gly Ser Ser Cys Gln Pro
                     1960
Cys Asp Cys Ser Gly Asn Gly Asp Pro Asn Leu Leu Phe Ser Asp Cys
   1970 1975
                                   1980
Asp Pro Leu Thr Gly Ala Cys Arg Gly Cys Leu Arg His Thr Thr Gly
    1990
                               1995
Pro Arg Cys Glu Ile Cys Ala Pro Gly Phe Tyr Gly Asn Ala Leu Leu
            2005
                            2010 2015
Pro Gly Asn Cys Thr Arg Cys Asp Cys Thr Pro Cys Gly Thr Glu Ala
        2020
                         2025
                                          2030
Cys Asp Pro His Ser Gly His Cys Leu Cys Lys Ala Gly Val Thr Gly
                      2040
                                      2045
Arg Arg Cys Asp Arg Cys Gln Glu Gly His Phe Gly Phe Asp Gly Cys
                  2055
                                   2060
Gly Gly Cys Arg Pro Cys Ala Cys Gly Pro Ala Ala Glu Gly Ser Glu
               2070
                                2075
Cys His Pro Gln Ser Gly Gln Cys His Cys Arg Pro Gly Thr Met Gly
            2085
                            2090
Pro Gln Cys Arg Glu Cys Ala Pro Gly Tyr Trp Gly Leu Pro Glu Gln
        2100
                                2110
                         2105
Gly Cys Arg Arg Cys Gln Cys Pro Gly Gly Arg Cys Asp Pro His Thr
                     2120
     2115
                                      2125
Gly Arg Cys Asn Cys Pro Pro Gly Leu Ser Gly Glu Arg Cys Asp Thr
  2130
                                   2140
                  2135
Cys Ser Gln Gln His Gln Val Pro Val Pro Gly Gly Pro Val Gly His
2145 2150 2155
Ser Ile His Cys Glu Val Cys Asp His Cys Val Val Leu Leu Asp
             2165
                    2170
```

	Leu	Glu	Arg 2180		Gly	Ala	Leu	Leu 2185		Ala	Ile	His			Leu
Arg	Gly	Ile 219	Asn		Ser	Ser	Met 2200	Ala		Ala	Arg	Leu 2205			Leu
Asn	Ala 2210	Ser		Ala	Asp	Leu 221	Gln		Gln	Leu	Arg 2220	Ser		Leu	Gly
Pro 222	Arg		Glu	Thr	Ala 2230	Gln		Leu	Glu	Val 2235	Leu		Gln	Gln	Ser 2240
Thr	Ser	Leu	Gly	Gln 2245		Ala	Arg	Arg	Leu 2250		Gly	Gln	Ala	Gly 2255	
	Arg		2260) -			_	2265	5			_	2270)	
	Leu	2275	5				2280)				2285	5		
Leu	Ala 2290		Ile	Arg	Ala	Val 2295		Arg	Thr	Leu	Ser 2300		Leu	Met	Ser
230		_			2310)				2315	5			-	2320
	Leu			2325	5				2330)				2335	5
_	Ala	_	2340)	_			2345	5				2350)	
	Ala	2355	5	_			2360)				2365	5		
	Trp 2370)				2375	5				2380)	=	•	
2385					2390)				2395	5				2400
	Val			2405	5				2410)			_	2415	5
	Arg		2420)				2425	,				2430)	
	Ala	2435	5				2440)				2445	5		
	Val 2450)				2455	5				2460)			
2465					2470)				2475	5				2480
	Met			Phe 2485		rro	Ala	GTÀ	Ser 2490		ьeu	Arg	ьeu	Val 2495	
АТА	АТА	ĢĽU		TT	70.7	\sim 1	~ 1	T -	α		T .	7.1	.		
0	0	T1 -	2500)				Leu 2505	,	Gln			2510	Asn)	Leu
	Ser	2515	2500 Ile) Leu	Asp	Val	Asn 2520	2505 Gln)	Asp	Gln Arg	Leu	Thr 2525	2510 Gln	Asn) Arg	Leu Ala
Ile	Glu 2530	2515 Ala)	2500 Ile Ser	Leu Asn	Asp Ala	Val Tyr 2535	Asn 2520 Ser	2505 Gln) Arg	Asp Ile	Gln Arg Leu	Leu Gln 2540	Thr 2525 Ala)	2510 Gln Val	Asn) Arg Gln	Leu Ala Ala
Ile Ala 2545	Glu 2530 Glu 5	2515 Ala) Asp	2500 Ile Ser Ala	Leu Asn Ala	Asp Ala Gly 2550	Val Tyr 2535 Gln	Asn 2520 Ser Ala	2505 Gln) Arg Leu	Asp Ile Gln	Gln Arg Leu Gln 2555	Leu Gln 2540 Ala	Thr 2525 Ala) Asp	2510 Gln Val His	Asn) Arg Gln Thr	Leu Ala Ala Trp 2560
Ile Ala 2545 Ala	Glu 2530 Glu 5 Thr	2515 Ala) Asp Val	2500 Ile Ser Ala Val	Leu Asn Ala Arg 2565	Asp Ala Gly 2550 Gln	Val Tyr 2535 Gln) Gly	Asn 2520 Ser Ala Leu	2505 Gln) Arg Leu Val	Asp Ile Gln Asp 2570	Gln Arg Leu Gln 2555 Arg	Leu Gln 2540 Ala Mla	Thr 2525 Ala) Asp Gln	2510 Gln Val His	Asn Arg Gln Thr Leu 2575	Leu Ala Ala Trp 2560 Leu
Ile Ala 2545 Ala Ala	Glu 2530 Glu 5 Thr Asn	2515 Ala) Asp Val	2500 Ile Ser Ala Val Thr 2580	Leu Asn Ala Arg 2565 Ala	Asp Ala Gly 2550 Gln Leu	Val Tyr 2535 Gln) Gly Glu	Asn 2520 Ser Ala Leu Glu	2505 Gln) Arg Leu Val Ala 2585	Asp Ile Gln Asp 2570 Met	Gln Arg Leu Gln 2555 Arg Leu	Leu Gln 2540 Ala Ala Gln	Thr 2525 Ala) Asp Gln	2510 Gln Val His Gln Gln 2590	Asn Arg Gln Thr Leu 2575 Gln	Leu Ala Ala Trp 2560 Leu Arg
Ile Ala 2545 Ala Ala Leu	Glu 2530 Glu 5 Thr	2515 Ala Asp Val Ser Leu 2595	2500 Ile Ser Ala Val Thr 2580 Val	Leu Asn Ala Arg 2565 Ala	Asp Ala Gly 2550 Gln Leu Ala	Val Tyr 2535 Gln) Gly Glu	Asn 2520 Ser Ala Leu Glu Leu 2600	2505 Gln Arg Leu Val Ala 2585 Gln	Asp Gln Asp 2570 Met Gly	Gln Arg Leu Gln 2555 Arg Leu Ala	Leu Gln 2540 Ala Ala Gln	Thr 2525 Ala) Asp Gln Glu Thr 2605	2510 Gln Val His Gln Gln 2590 Gln	Asn Arg Gln Thr Leu 2575 Gln Leu	Leu Ala Ala Trp 2560 Leu Arg

.

```
Gln Ala Met Leu Ala Met Asp Thr Asp Glu Thr Ser Lys Lys Ile Ala
         2630
                      2635
His Ala Lys Ala Val Ala Ala Glu Ala Gln Asp Thr Ala Thr Arg Val
             2645
                              2650
Gln Ser Gln Leu Gln Ala Met Gln Glu Asn Val Glu Arg Trp Gln Gly
         2660
                           2665
                                             2670
Gln Tyr Glu Gly Leu Arg Gly Gln Asp Leu Gly Gln Ala Val Leu Asp
                        2680
                               2685
Ala Gly His Ser Val Ser Thr Leu Glu Lys Thr Leu Pro Gln Leu Leu
                    2695
                                   2700
Ala Lys Leu Ser Ile Leu Glu Asn Arg Gly Val His Asn Ala Ser Leu
              2710
                                  2715
Ala Leu Ser Ala Ser Ile Gly Arg Val Arg Glu Leu Ile Ala Gln Ala
                               2730
              2725
Arg Gly Ala Ala Ser Lys Val Lys Val Pro Met Lys Phe Asn Gly Arg
                            2745
          2740
Ser Gly Val Gln Leu Arg Thr Pro Arg Asp Leu Ala Asp Leu Ala Ala
                        2760
                                          2765
Tyr Thr Ala Leu Lys Phe Tyr Leu Gln Gly Pro Glu Pro Glu Pro Gly
   2770 2775
                                       2780
Gln Gly Thr Glu Asp Arg Phe Val Met Tyr Met Gly Ser Arg Gln Ala
                2790
                                   2795
Thr Gly Asp Tyr Met Gly Val Ser Leu Arg Asp Lys Lys Val His Trp
              2805
                               2810
Val Tyr Gln Leu Gly Glu Ala Gly Pro Ala Val Leu Ser Ile Asp Glu
                            2825
Asp Ile Gly Glu Gln Phe Ala Ala Val Ser Leu Asp Arg Thr Leu Gln
                        2840
Phe Gly His Met Ser Val Thr Val Glu Arg Gln Met Ile Gln Glu Thr
                    2855
                                      2860
Lys Gly Asp Thr Val Ala Pro Gly Ala Glu Gly Leu Leu Asn Leu Arg
       2870
                                  2875
Pro Asp Asp Phe Val Phe Tyr Val Gly Gly Tyr Pro Ser Thr Phe Thr
                               2890
            2885
Pro Pro Pro Leu Leu Arg Phe Pro Gly Tyr Arg Gly Cys Ile Glu Met
         2900
                           2905
                                             2910
Asp Thr Leu Asn Glu Glu Val Val Ser Leu Tyr Asn Phe Glu Arg Thr
                        2920
                                          2925
Phe Gln Leu Asp Thr Ala Val Asp Arg Pro Cys Ala Arg Ser Lys Ser
                    2935
                                      2940
Thr Gly Asp Pro Trp Leu Thr Asp Gly Ser Tyr Leu Asp Gly Thr Gly
                2950
                                   2955
Phe Ala Arg Ile Ser Phe Asp Ser Gln Ile Ser Thr Thr Lys Arg Phe
             2965
                               2970
                                                 2975
Glu Gln Glu Leu Arg Leu Val Ser Tyr Ser Gly Val Leu Phe Phe Leu
          2980
                           2985
                                             2990
Lys Gln Gln Ser Gln Phe Leu Cys Leu Ala Val Gln Glu Gly Ser Leu
      2995 3000
                                          3005
Val Leu Leu Tyr Asp Phe Gly Ala Gly Leu Lys Lys Ala Val Pro Leu
                    3015
                                      3020
Gln Pro Pro Pro Leu Thr Ser Ala Ser Lys Ala Ile Gln Val Phe
                3030
                                   3035
Leu Leu Gly Gly Ser Arg Lys Arg Val Leu Val Arg Val Glu Arg Ala
             3045 3050
Thr Val Tyr Ser Val Glu Gln Asp Asn Asp Leu Glu Leu Ala Asp Ala
                           3065
```

```
Tyr Tyr Leu Gly Gly Val Pro Pro Asp Gln Leu Pro Pro Ser Leu Arg
 3075
              3080 3085
Arg Leu Phe Pro Thr Gly Gly Ser Val Arg Gly Cys Val Lys Gly Ile
                3095
                           3100
Lys Ala Leu Gly Lys Tyr Val Asp Leu Lys Arg Leu Asn Thr Thr Gly
      3110 3115
Val Ser Ala Gly Cys Thr Ala Asp Leu Leu Val Gly Arg Ala Met Thr
             3125
                   3130
Phe His Gly His Gly Phe Leu Arg Leu Ala Leu Ser Asn Val Ala Pro
                3145 3150
         3140
Leu Thr Gly Asn Val Tyr Ser Gly Phe Gly Phe His Ser Ala Gln Asp
      3155 3160 3165
Ser Ala Leu Leu Tyr Tyr Arg Ala Ser Pro Asp Gly Leu Cys Gln Val
                3175
                                    3180
Ser Leu Gln Gln Gly Arg Val Ser Leu Gln Leu Leu Arg Thr Glu Val
                3190
                                 3195
Lys Thr Gln Ala Gly Phe Ala Asp Gly Ala Pro His Tyr Val Ala Phe
             3205
                              3210
Tyr Ser Asn Ala Thr Gly Val Trp Leu Tyr Val Asp Asp Gln Leu Gln
                          3225
          3220
                                           3230
Gln Met Lys Pro His Arg Gly Pro Pro Pro Glu Leu Gln Pro Gln Pro
      3235
                       3240
                                        3245
Glu Gly Pro Pro Arg Leu Leu Gly Gly Leu Pro Glu Ser Gly Thr
                   3255
                                    3260
Ile Tyr Asn Phe Ser Gly Cys Ile Ser Asn Val Phe Val Gln Arg Leu
                3270
                                 3275
Leu Gly Pro Gln Arg Val Phe Asp Leu Gln Gln Asn Leu Gly Ser Val
                              3290
Asn Val Ser Thr Gly Cys Ala Pro Ala Leu Gln Ala Gln Thr Pro Gly
         3300
                          3305
Leu Gly Pro Arg Gly Leu Gln Ala Thr Ala Arg Lys Ala Ser Arg Arg
                      3320
                                       3325
Ser Arg Gln Pro Ala Arg His Pro Ala Cys Met Leu Pro Pro His Leu
                   3335
                                    3340
Arg Thr Thr Arg Asp Ser Tyr Gln Phe Gly Gly Ser Leu Ser Ser His
                                3355 3360
               3350
Leu Glu Phe Val Gly Ile Leu Ala Arg His Arg Asn Trp Pro Ser Leu
                             3370
             3365
                                              3375
Ser Met His Val Leu Pro Arg Ser Ser Arg Gly Leu Leu Leu Phe Thr
         3380
                          3385
                                           3390
Ala Arg Leu Arg Pro Gly Ser Pro Ser Leu Ala Leu Phe Leu Ser Asn
                       3400
                                        3405
Gly His Phe Val Ala Gln Met Glu Gly Leu Gly Thr Arg Leu Arg Ala
   3410
                   3415
                                    3420 -
Gln Ser Arg Gln Arg Ser Arg Pro Gly Arg Trp His Lys Val Ser Val
                3430
                                 3435
Arg Trp Glu Lys Asn Arg Ile Leu Leu Val Thr Asp Gly Ala Arg Ala
            3445
                             3450
                                              3455
Trp Ser Gln Glu Gly Pro His Arg Gln His Gln Gly Ala Glu His Pro
          3460
                          3465
                                           3470
Gln Pro His Thr Leu Phe Val Gly Gly Leu Pro Ala Ser Ser His Ser
     3475
                      3480 3485
Ser Lys Leu Pro Val Thr Val Gly Phe Ser Gly Cys Val Lys Arg Leu
  3490 3495
                                    3500
Arg Leu His Gly Arg Pro Leu Gly Ala Pro Thr Arg Met Ala Gly Val
                3510
                                 3515
```

Thr Pro Cys Ile Leu Gly Pro Leu Glu Ala Gly Leu Phe Phe Pro Gly 3525 3530 Ser Gly Gly Val Ile Thr Leu Asp Leu Pro Gly Ala Thr Leu Pro Asp 3545 3540 Val Gly Leu Glu Leu Glu Val Arg Pro Leu Ala Val Thr Gly Leu Ile 3565 3560 Phe His Leu Gly Gln Ala Arg Thr Pro Pro Tyr Leu Gln Leu Gln Val 3575 3580 Thr Glu Lys Gln Val Leu Leu Arg Ala Asp Asp Gly Ala Gly Glu Phe 3590 3595 Ser Thr Ser Val Thr Arg Pro Ser Val Leu Cys Asp Gly Gln Trp His 3610 Arg Leu Ala Val Met Lys Ser Gly Asn Val Leu Arg Leu Glu Val Asp 3625 Ala Gln Ser Asn His Thr Val Gly Pro Leu Leu Ala Ala Ala Gly 3640 3645 Ala Pro Ala Pro Leu Tyr Leu Gly Gly Leu Pro Glu Pro Met Ala Val 3655 3660 Gln Pro Trp Pro Pro Ala Tyr Cys Gly Cys Met Arg Arg Leu Ala Val 3670 3675 Asn Arg Ser Pro Val Ala Met Thr Arg Ser Val Glu Val His Gly Ala 3685 3690 Val Gly Ala Ser Gly Cys Pro Ala Ala

<210> 31 <211> 3696 <212> PRT <213> Homo sapiens

<400> 31

Met Ala Lys Arg Leu Cys Ala Gly Ser Ala Leu Cys Val Arg Gly Pro 10 Arg Gly Pro Ala Pro Leu Leu Val Gly Leu Ala Leu Leu Gly Ala 25 Ala Arg Ala Arg Glu Glu Ala Gly Gly Phe Ser Leu His Pro Pro 40 Tyr Phe Asn Leu Ala Glu Gly Ala Arg Ile Ala Ala Ser Ala Thr Cys Gly Glu Glu Ala Pro Ala Arg Gly Ser Pro Arg Pro Thr Glu Asp Leu 70 Tyr Cys Lys Leu Val Gly Gly Pro Val Ala Gly Gly Asp Pro Asn Gln 90 Thr Ile Arg Gly Gln Tyr Cys Asp Ile Cys Thr Ala Ala Asn Ser Asn 100 105 Lys Ala His Pro Ala Ser Asn Ala Ile Asp Gly Thr Glu Arg Trp Trp 120 115 125 Gln Ser Pro Pro Leu Ser Arg Gly Leu Glu Tyr Asn Glu Val Asn Val 135 140 Thr Leu Asp Leu Gly Gln Val Phe His Val Ala Tyr Val Leu Ile Lys 150 155 Phe Ala Asn Ser Pro Arg Pro Asp Leu Trp Val Leu Glu Arg Ser Met 170 Asp Phe Gly Arg Thr Tyr Gln Pro Trp Gln Phe Phe Ala Ser Ser Lys 185

```
Arg Asp Cys Leu Glu Arg Phe Gly Pro Gln Thr Leu Glu Arg Ile Thr
       195
                           200
Arg Asp Asp Ala Ala Ile Cys Thr Thr Glu Tyr Ser Arg Ile Val Pro
                        215
                                            220
Leu Glu Asn Gly Glu Ile Val Val Ser Leu Val Asn Gly Arg Pro Gly
                    230
                                        235
Ala Met Asn Phe Ser Tyr Ser Pro Leu Leu Arg Glu Phe Thr Lys Ala
                245
                                    250
Thr Asn Val Arg Leu Arg Phe Leu Arg Thr Asn Thr Leu Leu Gly His
                                265
Leu Met Gly Lys Ala Leu Arg Asp Pro Thr Val Thr Arg Arg Tyr Tyr
                            280
                                                285
Tyr Ser Ile Lys Asp Ile Ser Ile Gly Gly Arg Cys Val Cys His Gly
                        295
His Ala Asp Ala Cys Asp Ala Lys Asp Pro Thr Asp Pro Phe Arg Leu
                    310
                                        315
Gln Cys Thr Cys Gln His Asn Thr Cys Gly Gly Thr Cys Asp Arg Cys
                325
                                    330
Cys Pro Gly Phe Asn Gln Gln Pro Trp Lys Pro Ala Thr Ala Asn Ser
                                345
            340
Ala Asn Glu Cys Gln Ser Cys Asn Cys Tyr Gly His Ala Thr Asp Cys
                            360
                                                365
Tyr Tyr Asp Pro Glu Val Asp Arg Arg Ala Ser Gln Ser Leu Asp
                        375
                                            380
Gly Thr Tyr Gln Gly Gly Val Cys Ile Asp Cys Gln His His Thr
                    390
                                        395
Thr Gly Val Asn Cys Glu Arg Cys Leu Pro Gly Phe Tyr Arg Ser Pro
                405
                                    410
Asn His Pro Leu Asp Ser Pro His Val Cys Arg Arg Cys Asn Cys Glu
            420
                                425
Ser Asp Phe Thr Asp Gly Thr Cys Glu Asp Leu Thr Gly Arg Cys Tyr
                           440
                                                445
Cys Arg Pro Asn Phe Ser Gly Glu Arg Cys Asp Val Cys Ala Glu Gly
                       455
                                            460
Phe Thr Gly Phe Pro Ser Cys Tyr Pro Thr Pro Ser Ser Ser Asn Asp
                   .470
                                        475
Thr Arg Glu Gln Val Leu Pro Ala Gly Gln Ile Val Asn Cys Asp Cys
                485
                                    490
                                                        495
Ser Ala Ala Gly Thr Gln Gly Asn Ala Cys Arg Lys Asp Pro Arg Val
                                505
Gly Arg Cys Leu Cys Lys Pro Asn Phe Gln Gly Thr His Cys Glu Leu
                            520
Cys Ala Pro Gly Phe Tyr Gly Pro Gly Cys Gln Pro Cys Gln Cys Ser
                       535
                                            540
Ser Pro Gly Val Ala Asp Asp Arg Cys Asp Pro Asp Thr Gly Gln Cys
                   550
                                        555
Arg Cys Arg Val Gly Phe Glu Gly Ala Thr Cys Asp Arg Cys Ala Pro
                565
                                    570
Gly Tyr Phe His Phe Pro Leu Cys Gln Leu Cys Gly Cys Ser Pro Ala
           580
                                585
Gly Thr Leu Pro Glu Gly Cys Asp Glu Ala Gly Arg Cys Leu Cys Gln
                           600
Pro Glu Phe Ala Gly Pro His Cys Asp Arg Cys Arg Pro Gly Tyr His
                       615
                                            620
Gly Phe Pro Asn Cys Gln Ala Cys Thr Cys Asp Pro Arg Gly Ala Leu
```

```
Asp Gln Leu Cys Gly Ala Gly Gly Leu Cys Arg Cys Arg Pro Gly Tyr
              645
                           650
Thr Gly Thr Ala Cys Gln Glu Cys Ser Pro Gly Phe His Gly Phe Pro
           660
                              665
Ser Cys Val Pro Cys His Cys Ser Ala Glu Gly Ser Leu His Ala Ala
                           680
                                              685
Cys Asp Pro Arg Ser Gly Gln Cys Ser Cys Arg Pro Arg Val Thr Gly
                       695
                                           700
Leu Arg Cys Asp Thr Cys Val Pro Gly Ala Tyr Asn Phe Pro Tyr Cys
                                    715
                710
Glu Ala Gly Ser Cys His Pro Ala Gly Leu Ala Pro Val Asp Pro Ala
               725
                                   730
Leu Pro Glu Ala Gln Val Pro Cys Met Cys Arg Ala His Val Glu Gly
                               745
Pro Ser Cys Asp Arg Cys Lys Pro Gly Phe Trp Gly Leu Ser Pro Ser
                           760
Asn Pro Glu Gly Cys Thr Arg Cys Ser Cys Asp Leu Arg Gly Thr Leu
                       775
                                           780
Gly Gly Val Ala Glu Cys Gln Pro Gly Thr Gly Gln Cys Phe Cys Lys
                   790
                                       795
Pro His Val Cys Gly Gln Ala Cys Ala Ser Cys Lys Asp Gly Phe Phe
               805
                                   810
Gly Leu Asp Gln Ala Asp Tyr Phe Gly Cys Arg Ser Cys Arg Cys Asp
                               825
Ile Gly Gly Ala Leu Gly Gln Ser Cys Glu Pro Arg Thr Gly Val Cys
                           840
Arg Cys Arg Pro Asn Thr Gln Gly Pro Thr Cys Ser Glu Pro Ala Arg
                       855
Asp His Tyr Leu Pro Asp Leu His His Leu Arg Leu Glu Leu Glu Glu
                   870
                                       875
Ala Ala Thr Pro Glu Gly His Ala Val Arg Phe Gly Phe Asn Pro Leu
               885
                                   890
Glu Phe Glu Asn Phe Ser Trp Arg Gly Tyr Ala Gln Met Ala Pro Val
           900
                               905
Gln Pro Arg Ile Val Ala Arg Leu Asn Leu Thr Ser Pro Asp Leu Phe
                           920
                                               925
Trp Leu Val Phe Arg Tyr Val Asn Arg Gly Ala Met Ser Val Ser Gly
                       935
                                           940
Arg Val Ser Val Arg Glu Glu Gly Arg Ser Ala Thr Cys Ala Asn Cys
                   950
                                       955
Thr Ala Gln Ser Gln Pro Val Ala Phe Pro Pro Ser Thr Glu Pro Ala
                                   970
Phe Ile Thr Val Pro Gln Arg Gly Phe Gly Glu Pro Phe Val Leu Asn
           980
                               985
                                                   990
Pro Gly Thr Trp Ala Leu Arg Val Glu Ala Glu Gly Val Leu Leu Asp
        995
                           1000
                                               1005
Tyr Val Val Leu Leu Pro Ser Ala Tyr Tyr Glu Ala Ala Leu Leu Gln
                       1015
                                           1020
Leu Arg Val Thr Glu Ala Cys Thr Tyr Arg Pro Ser Ala Gln Gln Ser
                   1030
                                       1035
Gly Asp Asn Cys Leu Leu Tyr Thr His Leu Pro Leu Asp Gly Phe Pro
               1045
                                   1050
Ser Ala Ala Gly Leu Glu Ala Leu Cys Arg Gln Asp Asn Ser Leu Pro
                              1065
Arg Pro Cys Pro Thr Glu Gln Leu Ser Pro Ser His Pro Pro Leu Ile
                           1080
```

```
Thr Cys Thr Gly Ser Asp Val Asp Val Gln Leu Gln Val Ala Val Pro
  1090
         1095
                         1100
Gln Pro Gly Arg Tyr Ala Leu Val Val Glu Tyr Ala Asn Glu Asp Ala
             1110
                       1115
Arg Gln Glu Val Gly Val Ala Val His Thr Pro Gln Arg Ala Pro Gln
           1125
                   1130
Gln Gly Leu Leu Ser Leu His Pro Cys Leu Tyr Ser Thr Leu Cys Arg
       1140
                1145 1150
Gly Thr Ala Arg Asp Thr Gln Asp His Leu Ala Val Phe His Leu Asp
            1160
                                      1165
Ser Glu Ala Ser Val Arg Leu Thr Ala Glu Gln Ala Arg Phe Phe Leu
                  1175
                                   1180
His Gly Val Thr Leu Val Pro Ile Glu Glu Phe Ser Pro Glu Phe Val
               1190
                                1195
Glu Pro Arg Val Ser Cys Ile Ser Ser His Gly Ala Phe Gly Pro Asn
            1205
                             1210
Ser Ala Ala Cys Leu Pro Ser Arg Phe Pro Lys Pro Pro Gln Pro Ile
         1220
                         1225
                                         1230
Ile Leu Arg Asp Cys Gln Val Ile Pro Leu Pro Pro Gly Leu Pro Leu
      1235 1240
                                      1245
Thr His Ala Gln Asp Leu Thr Pro Ala Met Ser Pro Ala Gly Pro Arg
   1250 1255
                                   1260
Pro Arg Pro Pro Thr Ala Val Asp Pro Asp Ala Glu Pro Thr Leu Leu
               1270
                               1275
Arg Glu Pro Gln Ala Thr Val Val Phe Thr Thr His Val Pro Thr Leu
            1285
                            1290
Gly Arg Tyr Ala Phe Leu Leu His Gly Tyr Gln Pro Ala His Pro Thr
               1305
        1300
Phe Pro Val Glu Val Leu Ile Asn Ala Gly Arg Val Trp Gln Gly His
          1320
                                      1325
     1315
Ala Asn Ala Ser Phe Cys Pro His Gly Tyr Gly Cys Arg Thr Leu Val
  1330 1335
                                  1340
Val Cys Glu Gly Gln Ala Leu Leu Asp Val Thr His Ser Glu Leu Thr
    1350 1355
Val Thr Val Arg Val Pro Lys Gly Arg Trp Leu Trp Leu Asp Tyr Val
                            1370
                                            1375
            1365
Leu Val Val Pro Glu Asn Val Tyr Ser Phe Gly Tyr Leu Arg Glu Glu
         1380
                         1385
                                         1390
Pro Leu Asp Lys Ser Tyr Asp Phe Ile Ser His Cys Ala Ala Gln Gly
          1400
                                      1405
Tyr His Ile Ser Pro Ser Ser Ser Leu Phe Cys Arg Asn Ala Ala
   1410 1415
                                   1420
Ala Ser Leu Ser Leu Phe Tyr Asn Asn Gly Ala Arg Pro Cys Gly Cys
    1430
                               1435
His Glu Val Gly Ala Thr Gly Pro Thr Cys Glu Pro Phe Gly Gly Gln
            1445
                            1450
                                            1455
Cys Pro Cys His Ala His Val Ile Gly Arg Asp Cys Ser Arg Cys Ala
        1460
                         1465 1470
Thr Gly Tyr Trp Gly Phe Pro Asn Cys Arg Pro Cys Asp Cys Gly Ala
                     1480 1485
Arg Leu Cys Asp Glu Leu Thr Gly Gln Cys Ile Cys Pro Pro Arg Thr
                  1495
                                   1500
Ile Pro Pro Asp Cys Leu Leu Cys Gln Pro Gln Thr Phe Gly Cys His
             1510 1515
Pro Leu Val Gly Cys Glu Glu Cys Asn Cys Ser Gly Pro Gly Ile Gln
            1525
                            1530
```

```
Glu Leu Thr Asp Pro Thr Cys Asp Thr Asp Ser Gly Gln Cys Lys Cys
   1540 1545 1550
Arg Pro Asn Val Thr Gly Arg Arg Cys Asp Thr Cys Ser Pro Gly Phe
                1560
                            1565
His Gly Tyr Pro Arg Cys Arg Pro Cys Asp Cys His Glu Ala Gly Thr
  1570 1575 1580 ·
Ala Pro Gly Val Cys Asp Pro Leu Thr Gly Gln Cys Tyr Cys Lys Glu
         1590 1595
Asn Val Gln Gly Pro Lys Cys Asp Gln Cys Ser Leu Gly Thr Phe Ser
            1605
                 1610 1615
Leu Asp Ala Ala Asn Pro Lys Gly Cys Thr Arg Cys Phe Cys Phe Gly
         1620 1625 1630
Ala Thr Glu Arg Cys Arg Ser Ser Ser Tyr Thr Arg Gln Glu Phe Val
                   1640
                                     1645
Asp Met Glu Gly Trp Val Leu Leu Ser Thr Asp Arg Gln Val Val Pro
                  1655
                                   1660
His Glu Arg Gln Pro Gly Thr Glu Met Leu Arg Ala Asp Leu Arg His
               1670
                               1675
Val Pro Glu Ala Val Pro Glu Ala Phe Pro Glu Leu Tyr Trp Gln Ala
            1685 1690
Pro Pro Ser Tyr Leu Gly Asp Arg Val Ser Ser Tyr Gly Gly Thr Leu
         1700
                         1705
                                         1710
Arg Tyr Glu Leu His Ser Glu Thr Gln Arg Gly Asp Val Phe Val Pro
      1715 1720
                                     1725
Met Glu Ser Arg Pro Asp Val Val Leu Gln Gly Asn Gln Met Ser Ile
   1730 1735
                                  1740
Thr Phe Leu Glu Pro Ala Tyr Pro Thr Pro Gly His Val His Arg Gly
               1750
                               1755
Gln Leu Gln Leu Val Glu Gly Asn Phe Arg His Thr Glu Thr Arg Asn
            1765
                            1770
Thr Val Ser Arg Glu Glu Leu Met Met Val Leu Ala Ser Leu Glu Gln
        1780
                        1785
Leu Gln Ile Arg Ala Leu Phe Ser Gln Ile Ser Ser Ala Val Phe Leu
 1795
          1800
                                     1805
Arg Arg Val Ala Leu Glu Val Ala Ser Pro Ala Gly Gln Gly Ala Leu
                                  1820
  1810 1815
Ala Ser Asn Val Glu Leu Cys Leu Cys Pro Ala Ser Tyr Arg Gly Asp
              1830
                               1835
Ser Cys Gln Glu Cys Ala Pro Gly Phe Tyr Arg Asp Val Lys Gly Leu
            1845
                            1850
Phe Leu Gly Arg Cys Val Pro Cys Gln Cys His Gly His Ser Asp Arg
         1860
                         1865
                                         1870
Cys Leu Pro Gly Ser Gly Val Cys Val Asp Cys Gln His Asn Thr Glu
                     1880
                                      1885
Gly Ala His Cys Glu Arg Cys Gln Ala Gly Phe Val Ser Ser Arg Asp
                  1895
                                  1900
Asp Pro Ser Ala Pro Cys Val Ser Cys Pro Cys Pro Leu Ser Val Pro
     1910
                               1915
Ser Asn Asn Phe Ala Glu Gly Cys Val Leu Arg Gly Gly Arg Thr Gln
           1925
                            1930
                                             1935
Cys Leu Cys Lys Pro Gly Tyr Ala Gly Ala Ser Cys Glu Arg Cys Ala
        1940
                        1945
                                        1950
Pro Gly Phe Phe Gly Asn Pro Leu Val Leu Gly Ser Ser Cys Gln Pro
                     1960
                                     1965
Cys Asp Cys Ser Gly Asn Gly Asp Pro Asn Leu Leu Phe Ser Asp Cys
                  1975
                                   1980
```

Asp P: 1985	ro Leu	Thr G	Gly Al		Arg	Gly	Суѕ	Leu 1995	_	His	Thr	Thr	Gly 2000
Pro A	rg Cys		Ile Cy 2005	s Ala	Pro	Gly	Phe 2010		Gly	Asn	Ala	Leu 2015	
Pro G	ly Asn	Cys T 2020	Thr Ar	g Cys	Asp	Cys 2025		Pro	Cys	Gly	Thr 2030		Ala
Cys A	sp Pro 203		Ser Gl	y His	Cys 2040		Cys	Lys	Ala	Gly 2045		Thr	Gly
_	rg Cys 050	Asp A	Arg Cy	s Gln 205		Gly	His	Phe	Gly 2060		Asp	Gly	Cys
_	ly Cys	Arg F	Pro Cy 20	s Ala		Gly	Pro	Ala 2075	Ala		Gly	Ser	Glu 2080
Cys H	is Pro		Ser Gl 2085	y Gln	Cys	His	Cys 2090	Arg		Gly	Thr	Met 2095	_
Pro G	ln Cys	Arg G 2100	Glu Cy	s Ala	Pro	Gly 210		Trp	Gly	Leu	Pro 2110		Gln
Gly C	ys Arg 211		Cys Gl	n Cys	Pro 2120		Gly	Arg	Cys	Asp 2125		His	Thr
_	rg Cys 130	Asn C	Cys Pr	Pro 213		Leu	Ser	Gly	Glu 2140	_	Cys	Asp	Thr
Cys So 2145	er Gln	Gln H	His Gl		Pro	Val	Pro	Gly 2155	_	Pro	Val	Gly	His 2160
Ser I	le His	_	Glu Va 2165	l Cys	Asp	His	Cys 2170		Val	Leu	Leu	Leu 2175	_
Asp L	eu Glu	Arg A 2180	Ala Gl	y Ala	Leu	Leu 218		Ala	Ile	His	Glu 2190		Leu
Arg G	ly Ile 219		Ala Se	r Ser	Met 2200		Trp	Ala	Arg	Leu 2205		Arg	Leu
	la Ser 210	Ile A	Ala As	Leu 221		Ser	Gln	Leu	Arg 2220		Pro	Leu	Gly
Pro A: 2225	rg His	Glu T	Thr Al		Gln	Leu	Glu	Val 2235		Glu	Gln	Gln	Ser 2240
Thr S	er Leu		Gln As 2245	o Ala	Arg	Arg	Leu 2250		Gly	Gln	Ala	Ala 2255	
Gly T	hr Arg	Asp G 2260	Gln Al	a Ser	Gln	Leu 2265		Ala	Gly	Thr	Glu 2270		Thr
Leu G	ly His 227		Lys Th	r Leu	Leu 2280		Ala	Ile	Arg	Ala 2285		Asp	Arg
	eu Ser 290	Glu I	Leu Me	Ser 229		Thr	Gly	His	Leu 2300	_	Leu	Ala	Asn
Ala So 2305	er Ala	Pro S	Ser Gl 23		Gln	Leu	Leu	Arg 2315		Leu	Ala	Glu	Val 2320
Glu A	rg Leu		rp Gl [.] 2325	ı Met	Arg	Ala	Arg 2330		Leu	Gly	Ala	Pro 2335	
Ala A	la Ala	Glu A 2340	Ala Gl	ı Leu	Ala	Ala 2345		Gln	Arg	Leu	Leu 2350		Arg
Val G	ln Glu 235		Leu Se	r Ser	Leu 2360	_	Glu	Glu	Asn	Gln 2365		Leu	Ala
	ln Thr 370	Arg A	Asp Ar	g Leu 237		Gln	His	Glu	Ala 2380		Leu	Met	Asp
Leu A: 2385	rg Glu	Ala I	Leu As: 23	_	Ala	Val	Asp	Ala 2399		Arg	Glu	Ala	Gln 2400
Glu L	eu Asn		Arg As: 2405	n Gln	Glu	Arg	Leu 241		Glu	Ala	Leu	Gln 241	_
Lys G	ln Glu	Leu S 2420	Ser Ar	g Asp	Asn	Ala 242		Leu	Gln	Ala	Thr 2430		His

	2435	5				2440)		_		2445	5		
Asp Gln 2450		Lys	Glu	Glu	Leu 2455		Arg	Leu	Ala	Ala 2460		Leu	Asp	Gly
Ala Arg 2465	Thr	Pro	Leu	Leu 2470		Arg	Met	Gln	Thr 2475		Ser	Pro	Ala	Gly 2480
Ser Lys	Leu	Arg	Leu 2485		Glu	Ala	Ala	Glu 2490		His	Ala	Gln	Gln 2495	_
Gly Gln		2500)				2505	5			_	2510)	
	2515	5				2520)				2525	5		
Ile Leu 2530					2535	5				2540)			
Gln Gln . 2545				2550)				2555	5		_		2560
Asp Arg			2565	5				2570)				2575	5
Met Leu		2580)				2585	5				2590)	
	2595	5			-	2600)	-		_	2605	5		
Glu Ala 2610					2615	5				2620)	-		-
Glu Thr 2625		_	_	2630)			-	2635	5				2640
Gln Asp			2645	5				2650)				2655	5
Asn Val		2660)		_		2665	5			_	2670)	
	2675	5				2680)				2685	5		
Lys Thr					2695	<u> </u>				2700)			
Gly Val				2710)				2715	5				2720
Arg Glu			2725	5		_	_	2730)		-		2735	5
Pro Met		2740)				2745	5			_	2750)	_
	2755	.				2760)			_	2765	5		
Gly Pro 2770					2775	ò				2780)			
Tyr Met 2785				2790)				2795	5				2800
Arg Asp			2805	,	_		_	2810)	_			2815	5
Ala Val		2820)			_	2825	5				2830)	
	2835	,				2840)				2845	5		
Arg Gln i 2850					2855	, ,				2860)			
Glu Gly 1 2865	Leu	Leu	Asn	Leu 2870		Pro	Asp	Asp	Phe 2875		Phe	Tyr	Val	Gly 2880

```
Gly Tyr Pro Ser Thr Phe Thr Pro Pro Pro Leu Leu Arg Phe Pro Gly
             2885
                   2890 2895
Tyr Arg Gly Cys Ile Glu Met Asp Thr Leu Asn Glu Glu Val Val Ser
                            2905
          2900
                                               2910
Leu Tyr Asn Phe Glu Arg Thr Phe Gln Leu Asp Thr Ala Val Asp Arg
       2915
                        2920 2925
Pro Cys Ala Arg Ser Lys Ser Thr Gly Asp Pro Trp Leu Thr Asp Gly
                    2935 2940
Ser Tyr Leu Asp Gly Thr Gly Phe Ala Arg Ile Ser Phe Asp Ser Gln
                 2950
                                    2955
Ile Ser Thr Thr Lys Arg Phe Glu Gln Glu Leu Arg Leu Val Ser Tyr
              2965
                                2970
Ser Gly Val Leu Phe Phe Leu Lys Gln Gln Ser Gln Phe Leu Cys Leu
          2980
                             2985
                                               2990
Ala Val Gln Glu Gly Ser Leu Val Leu Leu Tyr Asp Phe Gly Ala Gly
                         3000
                                           3005
Leu Lys Lys Ala Val Pro Leu Gln Pro Pro Pro Pro Leu Thr Ser Ala
                     3015
                                        3020
Ser Lys Ala Ile Gln Val Phe Leu Leu Gly Gly Ser Arg Lys Arg Val
                 3030
                                    3035
Leu Val Arg Val Glu Arg Ala Thr Val Tyr Ser Val Glu Gln Asp Asn
              3045
                                3050
Asp Leu Glu Leu Ala Asp Ala Tyr Tyr Leu Gly Gly Val Pro Pro Asp
          3060
                            3065
Gln Leu Pro Pro Ser Leu Arg Arg Leu Phe Pro Thr Gly Gly Ser Val
                       3080
                                           3085
Arg Gly Cys Val Lys Gly Ile Lys Ala Leu Gly Lys Tyr Val Asp Leu
   3090 3095
                                       3100
Lys Arg Leu Asn Thr Thr Gly Val Ser Ala Gly Cys Thr Ala Asp Leu
                                   3115
      3110
Leu Val Gly Arg Ala Met Thr Phe His Gly His Gly Phe Leu Arg Leu
              3125
                               3130
Ala Leu Ser Asn Val Ala Pro Leu Thr Gly Asn Val Tyr Ser Gly Phe
          3140
                3145
Gly Phe His Ser Ala Gln Asp Ser Ala Leu Leu Tyr Tyr Arg Ala Ser
                        3160
      3155
                                           3165
Pro Asp Gly Leu Cys Gln Val Ser Leu Gln Gln Gly Arg Val Ser Leu
                     3175
                                        3180
Gln Leu Leu Arg Thr Glu Val Lys Thr Gln Ala Gly Phe Ala Asp Gly
                 3190
                                    3195
Ala Pro His Tyr Val Ala Phe Tyr Ser Asn Ala Thr Gly Val Trp Leu
             3205
                                3210
                                                   3215
Tyr Val Asp Asp Gln Leu Gln Gln Met Lys Pro His Arg Gly Pro Pro
          3220
                            3225
                                               3230
Pro Glu Leu Gln Pro Gln Pro Glu Gly Pro Pro Arg Leu Leu Gly
      3235
                        3240
                                           3245
Gly Leu Pro Glu Ser Gly Thr Ile Tyr Asn Phe Ser Gly Cys Ile Ser
                     3255
                                        3260
Asn Val Phe Val Gln Arg Leu Leu Gly Pro Gln Arg Val Phe Asp Leu
                3270
                                   3275
Gln Gln Asn Leu Gly Ser Val Asn Val Ser Thr Gly Cys Ala Pro Ala
              3285
                                3290
Leu Gln Ala Gln Thr Pro Gly Leu Gly Pro Arg Gly Leu Gln Ala Thr
                            3305
Ala Arg Lys Ala Ser Arg Arg Ser Arg Gln Pro Ala Arg His Pro Ala
                        3320
```

```
Cys Met Leu Pro Pro His Leu Arg Thr Thr Arg Asp Ser Tyr Gln Phe
   3330
         3335
                          3340
Gly Gly Ser Leu Ser Ser His Leu Glu Phe Val Gly Ile Leu Ala Arg
          3350
3345
                                3355
His Arg Asn Trp Pro Ser Leu Ser Met His Val Leu Pro Arg Ser Ser
             3365
                    3370
Arg Gly Leu Leu Phe Thr Ala Arg Leu Arg Pro Gly Ser Pro Ser
                         3385
          3380
Leu Ala Leu Phe Leu Ser Asn Gly His Phe Val Ala Gln Met Glu Gly
                       3400
                                          3405
Leu Gly Thr Arg Leu Arg Ala Gln Ser Arg Gln Arg Ser Arg Pro Gly
                    3415
                                      3420
Arg Trp His Lys Val Ser Val Arg Trp Glu Lys Asn Arg Ile Leu Leu
3425
                 3430
                                   3435
Val Thr Asp Gly Ala Arg Ala Trp Ser Gln Glu Gly Pro His Arg Gln
              3445
                               3450
His Gln Gly Ala Glu His Pro Gln Pro His Thr Leu Phe Val Gly Gly
          3460
                           3465
                                             3470
Leu Pro Ala Ser Ser His Ser Ser Lys Leu Pro Val Thr Val Gly Phe
                       3480
                                          3485
Ser Gly Cys Val Lys Arg Leu Arg Leu His Gly Arg Pro Leu Gly Ala
                    3495
                                      3500
Pro Thr Arg Met Ala Gly Val Thr Pro Cys Ile Leu Gly Pro Leu Glu
                3510
                                  3515
Ala Gly Leu Phe Phe Pro Gly Ser Gly Gly Val Ile Thr Leu Asp Leu
             3525
                               3530
Pro Gly Ala Thr Leu Pro Asp Val Gly Leu Glu Leu Glu Val Arg Pro
          3540
                           3545
Leu Ala Val Thr Gly Leu Ile Phe His Leu Gly Gln Ala Arg Thr Pro
                        3560
                                         3565
Pro Tyr Leu Gln Leu Gln Val Thr Glu Lys Gln Val Leu Leu Arg Ala
   3570 3575
                                      3580
Asp Asp Gly Ala Gly Glu Phe Ser Thr Ser Val Thr Arg Pro Ser Val
                3590 . 3595
Leu Cys Asp Gly Gln Trp His Arg Leu Ala Val Met Lys Ser Gly Asn
             3605 3610
Val Leu Arg Leu Glu Val Asp Ala Gln Ser Asn His Thr Val Gly Pro
          3620
                           3625
Leu Leu Ala Ala Ala Gly Ala Pro Ala Pro Leu Tyr Leu Gly Gly
                       3640
                                         3645
Leu Pro Glu Pro Met Ala Val Gln Pro Trp Pro Pro Ala Tyr Cys Gly
                    3655
                                      3660
Cys Met Arg Arg Leu Ala Val Asn Arg Ser Pro Val Ala Met Thr Arg
3665 3670
                                  3675
Ser Val Glu Val His Gly Ala Val Gly Ala Ser Gly Cys Pro Ala Ala
              3685
                               3690
```

```
<210> 32
```

<400> 32

Met Thr Asn Asn Ser Gly Ser Lys Ala Glu Leu Val Val Gly Gly Lys

1 10 15

<211> 337

<212> PRT

<213> Homo sapiens

```
Tyr Lys Leu Val Arg Lys Ile Gly Ser Gly Ser Phe Gly Asp Val Tyr
            20
                                25
Leu Gly Ile Thr Thr Asn Gly Glu Asp Val Ala Val Lys Leu Glu
                            40
Ser Gln Lys Val Lys His Pro Gln Leu Leu Tyr Glu Ser Lys Leu Tyr
                        55
Thr Ile Leu Gln Gly Gly Val Gly Ile Pro His Met His Trp Tyr Gly
                    70
                                        75
Gln Glu Lys Asp Asn Asn Val Leu Val Met Asp Leu Leu Gly Pro Ser
                8.5
                                    90
Leu Glu Asp Leu Phe Asn Phe Cys Ser Arg Arg Phe Thr Met Lys Thr
                                105
Val Leu Met Leu Ala Asp Gln Met Ile Ser Arg Ile Glu Tyr Val His
                            120
Thr Lys Asn Phe Leu His Arg Asp Ile Lys Pro Asp Asn Phe Leu Met
    130
                        135
                                            140
Gly Thr Gly Arg His Cys Asn Lys Leu Phe Leu Ile Asp Phe Gly Leu
                    150
                                        155
Ala Lys Lys Tyr Arg Asp Asn Arg Thr Arg Gln His Ile Pro Tyr Arg
                165
                                    170
                                                        175
Glu Asp Lys His Leu Ile Gly Thr Val Arg Tyr Ala Ser Ile Asn Ala
            180
                                185
                                                    190
His Leu Gly Ile Glu Gln Ser Arg Arg Asp Asp Met Glu Ser Leu Gly
                            200
                                                205
Tyr Val Phe Met Tyr Phe Asn Arg Thr Ser Leu Pro Trp Gln Gly Leu
                        215
                                            220
Arg Ala Met Thr Lys Lys Gln Lys Tyr Glu Lys Ile Ser Glu Lys Lys
                    230
                                        235
Met Ser Thr Pro Val Glu Val Leu Cys Lys Gly Phe Pro Ala Glu Phe
                245
                                   250
Ala Met Tyr Leu Asn Tyr Cys Arg Gly Leu Arg Phe Glu Glu Val Pro
           260
                                265
Asp Tyr Met Tyr Leu Arg Gln Leu Phe Arg Ile Leu Phe Arg Thr Leu
                            280
                                                285
Asn His Gln Tyr Asp Tyr Thr Phe Asp Trp Thr Met Leu Lys Gln Lys
                        295
                                            300
Ala Ala Gln Gln Ala Ala Ser Ser Gly Gln Gly Gln Gln Ala Gln
                   310
                                       315
Thr Gln Thr Gly Lys Gln Thr Glu Lys Asn Lys Asn Asn Val Lys Asp
Asn
```

```
<210> 33
```

 Met Glu Ser Leu Leu Leu Leu Pro Val Leu Leu Leu Leu Ala Ile Leu Trp

 1
 5
 10
 15

 Thr Gln Ala Ala Ala Leu Ile Asn Leu Lys Tyr Ser Val Glu Glu Glu
 20
 25
 30

 Gln Arg Ala Gly Thr Val Ile Ala Asn Val Ala Lys Asp Ala Arg Glu
 45

<211> 888

<212> PRT

<213> Homo sapiens

<400> 33

```
Ala Gly Phe Ala Leu Asp Pro Arg Gln Ala Ser Ala Phe Arg Val Val
                       55
Ser Asn Ser Ala Pro His Leu Val Asp Ile Asn Pro Ser Ser Gly Leu
                                        7.5
Leu Val Thr Lys Gln Lys Ile Asp Arg Asp Leu Leu Cys Arg Gln Ser
                                    90
                85
Pro Lys Cys Ile Ile Ser Leu Glu Val Met Ser Ser Met Glu Ile
                                105
Cys Val Ile Lys Val Glu Ile Lys Asp Leu Asn Asp Asn Ala Pro Ser
                            120
Phe Pro Ala Ala Gln Ile Glu Leu Glu Ile Ser Glu Ala Ala Ser Pro
                        135
Gly Thr Arg Ile Pro Leu Asp Ser Ala Tyr Asp Pro Asp Ser Gly Ser
                    150
                                        155
Phe Gly Val Gln Thr Tyr Glu Leu Thr Pro Asn Glu Leu Phe Gly Leu
                165
                                    170
                                                        175
Glu Ile Lys Thr Arg Gly Asp Gly Ser Arg Phe Ala Glu Leu Val Val
           180
                                185
Glu Lys Ser Leu Asp Arg Glu Thr Gln Ser His Tyr Ser Phe Arg Ile
                            200
                                                205
Thr Ala Leu Asp Gly Gly Asp Pro Pro Arg Leu Gly Thr Val Gly Leu
                       215
Ser Ile Lys Val Thr Asp Ser Asn Asp Asn Asn Pro Val Phe Ser Glu
                    230
                                        235
Ser Thr Tyr Ala Val Ser Val Pro Glu Asn Ser Pro Pro Asn Thr Pro
               245
                                    250
Val Ile Arg Leu Asn Ala Ser Asp Pro Asp Glu Gly Thr Asn Gly Gln
           260
                               265
Val Val Tyr Ser Phe Tyr Gly Tyr Val Asn Asp Arg Thr Arg Glu Leu
                            280
                                                285
Phe Gln Ile Asp Pro His Ser Gly Leu Val Thr Val Thr Gly Ala Leu
                       295
                                            300
Asp Tyr Glu Glu Gly His Val Tyr Glu Leu Asp Val Gln Ala Lys Asp
                    310
                                       315
Leu Gly Pro Asn Ser Ile Pro Ala His Cys Lys Val Thr Val Ser Val
                325
                                    330
Leu Asp Thr Asn Asp Asn Pro Pro Val Ile Asn Leu Leu Ser Val Asn
                                345
Ser Glu Leu Val Glu Val Ser Glu Ser Ala Pro Pro Gly Tyr Val Ile
                            360
Ala Leu Val Arg Val Ser Asp Arg Asp Ser Gly Leu Asn Gly Arg Val
                        375
                                            380
Gln Cys Arg Leu Leu Gly Asn Val Pro Phe Arg Leu Gln Glu Tyr Glu
                    390
                                        395
Ser Phe Ser Thr Ile Leu Val Asp Gly Arg Leu Asp Arg Glu Gln His
               405
                                   410
Asp Gln Tyr Asn Leu Thr Ile Gln Ala Arg Asp Gly Gly Val Pro Met
           420
                               425
Leu Gln Ser Ala Lys Ser Phe Thr Val Leu Ile Thr Asp Glu Asn Asp
                           440
                                                445
Asn His Pro His Phe Ser Lys Pro Tyr Tyr Gln Val Ile Val Gln Glu
                       455
                                            460
Asn Asn Thr Pro Gly Ala Tyr Leu Leu Ser Val Ser Ala Arg Asp Pro
                   470
                                       475
Asp Leu Gly Leu Asn Gly Ser Val Ser Tyr Gln Ile Val Pro Ser Gln
                                    490
```

```
Val Arg Asp Met Pro Val Phe Thr Tyr Val Ser Ile Asn Pro Asn Ser
          500
                    505
Gly Asp Ile Tyr Ala Leu Arg Ser Phe Asn His Glu Gln Thr Lys Ala
                           520
                                               525
Phe Glu Phe Lys Val Leu Ala Lys Asp Gly Gly Leu Pro Ser Leu Gln
                       535
                                           540
Ser Asn Ala Thr Val Arg Val Ile Ile Leu Asp Val Asn Asp Asn Thr
                   550
                                       555
Pro Val Ile Thr Ala Pro Pro Leu Ile Asn Gly Thr Ala Glu Val Tyr
               565
                                   570
Ile Pro Arg Asn Ser Gly Ile Gly Tyr Leu Val Thr Val Val Lys Ala
                               585
Glu Asp Tyr Asp Glu Gly Glu Asn Gly Arg Val Thr Tyr Asp Met Thr
                           600
                                               605
Glu Gly Asp Arg Gly Phe Phe Glu Ile Asp Gln Val Asn Gly Glu Val
                       615
                                           620
Arg Thr Thr Arg Thr Phe Gly Glu Ser Ser Lys Ser Ser Tyr Glu Leu
                   630
                                       635
Ile Val Val Ala His Asp His Gly Lys Thr Ser Leu Ser Ala Ser Ala
               645
                                   650
Leu Val Leu Ile Tyr Leu Ser Pro Ala Leu Asp Ala Gln Glu Ser Met
           660
                               665
Gly Ser Val Asn Leu Ser Leu Ile Phe Ile Ile Ala Leu Gly Ser Ile
                           680
Ala Gly Ile Leu Phe Val Thr Met Ile Phe Val Ala Ile Lys Cys Lys
                       695
Arg Asp Asn Lys Glu Ile Arg Thr Tyr Asn Cys Ser Asn Cys Leu Thr
                  710
                                       715
Ile Thr Cys Leu Leu Gly Cys Phe Ile Lys Gly Gln Asn Ser Lys Cys
               725
                                  730
Leu His Cys Ile Ser Val Ser Pro Ile Ser Glu Glu Gln Asp Lys
                              745
           740
                                                   750
Thr Glu Glu Lys Val Ser Leu Arg Gly Lys Arg Ile Ala Glu Tyr Ser
                          760
                                               765
Tyr Gly His Gln Lys Lys Ser Ser Lys Lys Lys Ile Ser Lys Asn
                      775
                                           780
Asp Ile Arg Leu Val Pro Arg Asp Val Glu Glu Thr Asp Lys Met Asn
                   790
                                       795
Val Val Ser Cys Ser Ser Leu Thr Ser Ser Leu Asn Tyr Phe Asp Tyr
               805
                                   810
His Gln Gln Thr Leu Pro Leu Gly Cys Arg Arg Ser Glu Ser Thr Phe
                               825
Leu Asn Val Glu Asn Gln Asn Thr Arg Asn Thr Ser Ala Asn His Ile
                           840
Tyr His His Ser Phe Asn Ser Gln Gly Pro Gln Gln Pro Asp Leu Ile
                      855
                                           860
Ile Asn Gly Val Pro Leu Pro Glu Val Ser Ala Ala Lys Trp Leu Cys
                870
                                      875
Glu Val Leu Pro Gly Leu Leu Leu
               885
```

<210> 34 <211> 855

<211> 833 <212> PRT

<213> Homo sapiens

<400> 34 Met Glu Ser Leu Leu Pro Val Leu Leu Leu Ala Ile Leu Trp Thr Gln Ala Ala Leu Ile Asn Leu Lys Tyr Ser Val Glu Glu Glu Gln Arg Ala Gly Thr Val Ile Ala Asn Val Ala Lys Asp Ala Arg Glu Ala Gly Phe Ala Leu Asp Pro Arg Gln Ala Ser Ala Phe Arg Val Val · 55 Ser Asn Ser Ala Pro His Leu Val Asp Ile Asn Pro Ser Ser Gly Leu Leu Val Thr Lys Gln Lys Ile Asp Arg Asp Leu Leu Cys Arg Gln Ser Pro Lys Cys Ile Ile Ser Leu Glu Val Met Ser Ser Met Glu Ile Cys Val Ile Lys Val Glu Ile Lys Asp Leu Asn Asp Asn Ala Pro Ser Phe Pro Ala Ala Gln Ile Glu Leu Glu Ile Ser Glu Ala Ala Ser Pro Gly Thr Arg Ile Pro Leu Asp Ser Ala Tyr Asp Pro Asp Ser Gly Ser Phe Gly Val Gln Thr Tyr Glu Leu Thr Pro Asn Glu Leu Phe Gly Leu Glu Ile Lys Thr Arg Gly Asp Gly Ser Arg Phe Ala Glu Leu Val Val Glu Lys Ser Leu Asp Arg Glu Thr Gln Ser His Tyr Ser Phe Arg Ile Thr Ala Leu Asp Gly Gly Asp Pro Pro Arg Leu Gly Thr Val Gly Leu Ser Ile Lys Val Thr Asp Ser Asn Asp Asn Asn Pro Val Phe Ser Glu Ser Thr Tyr Ala Val Ser Val Pro Glu Asn Ser Pro Pro Asn Thr Pro Val Ile Arg Leu Asn Ala Ser Asp Pro Asp Glu Gly Thr Asn Gly Gln Val Val Tyr Ser Phe Tyr Gly Tyr Val Asn Asp Arg Thr Arg Glu Leu Phe Gln Ile Asp Pro His Ser Gly Leu Val Thr Val Thr Gly Ala Leu Asp Tyr Glu Glu Gly His Val Tyr Glu Leu Asp Val Gln Ala Lys Asp Leu Gly Pro Asn Ser Ile Pro Ala His Cys Lys Val Thr Val Ser Val Leu Asp Thr Asn Asp Asn Pro Pro Val Ile Asn Leu Leu Ser Val Asn Ser Glu Leu Val Glu Val Ser Glu Ser Ala Pro Pro Gly Tyr Val Ile Ala Leu Val Arg Val Ser Asp Arg Asp Ser Gly Leu Asn Gly Arg Val Gln Cys Arg Leu Leu Gly Asn Val Pro Phe Arg Leu Gln Glu Tyr Glu Ser Phe Ser Thr Ile Leu Val Asp Gly Arg Leu Asp Arg Glu Gln His Asp Gln Tyr Asn Leu Thr Ile Gln Ala Arg Asp Gly Gly Val Pro Met Leu Gln Ser Ala Lys Ser Phe Thr Val Leu Ile Thr Asp Glu Asn Asp

```
Asn His Pro His Phe Ser Lys Pro Tyr Tyr Gln Val Ile Val Gln Glu
                     455
Asn Asn Thr Pro Gly Ala Tyr Leu Leu Ser Val Ser Ala Arg Asp Pro
                 470
                                      475
Asp Leu Gly Leu Asn Gly Ser Val Ser Tyr Gln Ile Val Pro Ser Gln
              485
                       490
Val Arg Asp Met Pro Val Phe Thr Tyr Val Ser Ile Asn Pro Asn Ser
          500
                             505
Gly Asp Ile Tyr Ala Leu Arg Ser Phe Asn His Glu Gln Thr Lys Ala
                          520
                                             525
Phe Glu Phe Lys Val Leu Ala Lys Asp Gly Gly Leu Pro Ser Leu Gln
                       535
                                          540
Ser Asn Ala Thr Val Arg Val Ile Ile Leu Asp Val Asn Asp Asn Thr
                   550
                                      555
Pro Val Ile Thr Ala Pro Pro Leu Ile Asn Gly Thr Ala Glu Val Tyr
                                   570
Ile Pro Arg Asn Ser Gly Ile Gly Tyr Leu Val Thr Val Val Lys Ala
                               585
Glu Asp Tyr Asp Glu Gly Glu Asn Gly Arg Val Thr Tyr Asp Met Thr
       595
                           600
                                               605
Glu Gly Asp Arg Gly Phe Phe Glu Ile Asp Gln Val Asn Gly Glu Val
                       615
                                          620
Arg Thr Thr Arg Thr Phe Gly Glu Ser Ser Lys Ser Ser Tyr Glu Leu
                   630
                                      635
Ile Val Val Ala His Asp His Gly Lys Thr Ser Leu Ser Ala Ser Ala
                                  650
Leu Val Leu Ile Tyr Leu Ser Pro Ala Leu Asp Ala Gln Glu Ser Met
                               665
Gly Ser Val Asn Leu Ser Leu Ile Phe Ile Ile Ala Leu Gly Ser Ile
                          680
Ala Gly Ile Leu Phe Val Thr Met Ile Phe Val Ala Ile Lys Cys Lys
                       695
                                          700
Arg Asp Asn Lys Glu Ile Arg Thr Tyr Asn Cys Arg Ile Ala Glu Tyr
                  710
                                      715
Ser Tyr Gly His Gln Lys Lys Ser Ser Lys Lys Lys Ile Ser Lys
               725
                                 730
Asn Asp Ile Arg Leu Val Pro Arg Asp Val Glu Glu Thr Asp Lys Met
           740
                              745
                                                  750
Asn Val Val Ser Cys Ser Ser Leu Thr Ser Ser Leu Asn Tyr Phe Asp
                           760
                                   .
Tyr His Gln Gln Thr Leu Pro Leu Gly Cys Arg Arg Ser Glu Ser Thr
                       775
Phe Leu Asn Val Glu Asn Gln Asn Thr Arg Asn Thr Ser Ala Asn His
                   790
                                      795
Ile Tyr His His Ser Phe Asn Ser Gln Gly Pro Gln Gln Pro Asp Leu
              805
                                  810
Ile Ile Asn Gly Val Pro Leu Pro Glu Thr Glu Asn Tyr Ser Phe Asp
               825
        820
                                                 830
Ser Asn Tyr Val Asn Ser Arg Ala His Leu Ile Lys Arg Tyr Val Gly
              840
Leu Leu Ala Tyr Cys Cys Asn
```

<210> 35 <211> 329

<212> PRT

<213> Homo sapiens

```
<400> 35
Met Val Thr Lys Ala Phe Val Leu Leu Ala Ile Phe Ala Glu Ala Ser
                                   10
Ala Lys Ser Cys Ala Pro Asn Lys Ala Asp Val Ile Leu Val Phe Cys
        20
                               25
Tyr Pro Lys Thr Ile Ile Thr Lys Ile Pro Glu Cys Pro Tyr Gly Trp
                           40
Glu Val His Gln Leu Ala Leu Gly Gly Leu Cys Tyr Asn Gly Val His
                        55
                                           60
Glu Gly Gly Tyr Tyr Gln Phe Val Ile Pro Asp Leu Ser Pro Lys Asn
                                        75
Lys Ser Tyr Cys Gly Thr Gln Ser Glu Tyr Lys Pro Pro Ile Tyr His
                                    90
Phe Tyr Ser His Ile Val Ser Asn Asp Thr Thr Val Ile Val Lys Asn
            100
                                105
Gln Pro Val Asn Tyr Ser Phe Ser Cys Thr Tyr His Ser Thr Tyr Leu
                            120
Val Asn Gln Ala Ala Phe Asp Gln Arg Val Ala Thr Val His Val Lys
                        135
                                            140
Asn Gly Ser Met Gly Thr Phe Glu Ser Gln Leu Ser Leu Asn Phe Tyr
                    150
                                        155
Thr Asn Ala Lys Phe Ser Ile Lys Lys Glu Ala Pro Phe Val Leu Glu
               165
                                    170
Ala Ser Glu Ile Gly Ser Asp Leu Phe Ala Gly Val Glu Ala Lys Gly
                                185
Leu Ser Ile Arg Phe Lys Val Val Leu Asn Ser Cys Trp Ala Thr Pro
                           200
Ser Ala Asp Phe Met Tyr Pro Leu Gln Trp Gln Leu Ile Asn Lys Gly
                       215
                                            220
Cys Pro Thr Asp Glu Thr Val Leu Val His Glu Asn Gly Arg Asp His
                   230
                                        235
Arg Ala Thr Phe Gln Phe Asn Ala Phe Arg Phe Gln Asn Ile Pro Lys
               245
                                    250
Leu Ser Lys Val Trp Leu His Cys Glu Thr Phe Ile Cys Asp Ser Glu
           260
                               265
Lys Leu Ser Cys Pro Val Thr Cys Asp Lys Arg Lys Arg Leu Leu Arg
                           280
                                               285
Asp Gln Thr Gly Gly Val Leu Val Val Glu Leu Ser Leu Arg Ser Arg
                        295
                                            300
Gly Phe Ser Ser Leu Tyr Ser Phe Ser Asp Val Leu His His Leu Ile
                    310
                                       315
Met Met Leu Gly Ile Cys Ala Val Leu
               325
```

```
<210> 36
<211> 232
<212> PRT
<213> Homo sapiens
```

<400> 36

Met Leu Tyr Thr Arg Lys Asn Leu Thr Cys Ala Gln Thr Ile Asn Ser 1 5 10 15

Ser Ala Phe Gly Asn Leu Asn Val Thr Lys Lys Thr Thr Phe Ile Val 20 25 30

```
His Gly Phe Arg Pro Thr Gly Ser Pro Pro Val Trp Met Asp Asp Leu
                           40
Val Lys Gly Leu Leu Ser Val Glu Asp Met Asn Val Val Val Asp
                       55
Trp Asn Arg Gly Ala Thr Thr Leu Ile Tyr Thr His Ala Ser Ser Lys
                   70
                                       75
Thr Arg Lys Val Ala Met Val Leu Lys Glu Phe Ile Asp Gln Met Leu
                                    90
Ala Glu Gly Ala Ser Leu Asp Asp Ile Tyr Met Ile Gly Val Ser Leu
           100
                               105
Gly Ala His Ile Ser Gly Phe Val Gly Glu Met Tyr Asp Gly Trp Leu
                            120
                                                125
Gly Arg Ile Thr Gly Leu Asp Pro Ala Gly Pro Leu Phe Asn Gly Lys
                        135
                                            140
Pro His Gln Asp Arg Leu Asp Pro Ser Asp Ala Gln Phe Val Asp Val
                                        155
Ile His Ser Asp Thr Asp Gly Asn Ala Pro Phe Leu Val Ala Leu Gly
                165
                                    170
Tyr Lys Glu Pro Leu Gly Asn Ile Asp Phe Tyr Pro Asn Gly Gly Leu
                                185
           180
Asp Gln Pro Gly Cys Pro Lys Thr Ile Leu Gly Gly Asn Val Lys Glu
                            200
                                                205
Met Ile Gln Ala Ser Tyr Ile Phe Phe Leu Lys Asn Asp Ser Met Asp
                        215
Leu Ser Ser Pro Lys Glu Val Glu
```

<211> 452

<212> PRT

<213> Homo sapiens

<400> 37

Met Leu Arg Phe Tyr Leu Phe Ile Ser Leu Leu Cys Leu Ser Arg Ser 10 Asp Ala Glu Glu Thr Cys Pro Ser Phe Thr Arg Leu Ser Phe His Ser 25 Ala Val Val Gly Thr Gly Leu Asn Val Arg Leu Met Leu Tyr Thr Arg 40 Lys Asn Leu Thr Cys Ala Gln Thr Ile Asn Ser Ser Ala Phe Gly Asn Leu Asn Val Thr Lys Lys Thr Thr Phe Ile Val His Gly Phe Arg Pro 70 7.5 Thr Gly Ser Pro Pro Val Trp Met Asp Asp Leu Val Lys Gly Leu Leu 85 90 Ser Val Glu Asp Met Asn Val Val Val Asp Trp Asn Arg Gly Ala 100 105 Thr Thr Leu Ile Tyr Thr His Ala Ser Ser Lys Thr Arg Lys Val Ala 120 125 Met Val Leu Lys Glu Phe Ile Asp Gln Met Leu Ala Glu Gly Ala Ser 135 140 Leu Asp Asp Ile Tyr Met Ile Gly Val Ser Leu Gly Ala His Ile Ser 150 155 Gly Phe Val Gly Glu Met Tyr Asp Gly Trp Leu Gly Arg Ile Thr Gly

```
Leu Asp Pro Ala Gly Pro Leu Phe Asn Gly Lys Pro His Gln Asp Arg
           180
                               185
Leu Asp Pro Ser Asp Ala Gln Phe Val Asp Val Ile His Ser Asp Thr
                            200
Asp Ala Leu Gly Tyr Lys Glu Pro Leu Gly Asn Ile Asp Phe Tyr Pro
                        215
                                            220
Asn Gly Gly Leu Asp Gln Pro Gly Cys Pro Lys Thr Ile Leu Gly Gly
                    230
                                        235
Phe Gln Tyr Phe Lys Cys Asp His Gln Arg Ser Val Tyr Leu Tyr Leu
                245
                                    250
Ser Ser Leu Arg Glu Ser Cys Thr Ile Thr Ala Tyr Pro Cys Asp Ser
                                265
                                                    270
Tyr Gln Asp Tyr Arg Asn Gly Lys Cys Val Ser Cys Gly Thr Ser Gln
                            280
Lys Glu Ser Cys Pro Leu Leu Gly Tyr Tyr Ala Asp Asn Trp Lys Asp
                        295
                                            300
His Leu Arg Gly Lys Asp Pro Pro Met Thr Lys Ala Phe Phe Asp Thr
                    310
                                        315
Ala Glu Glu Ser Pro Phe Cys Met Tyr His Tyr Phe Val Asp Ile Ile
                325
                                    330
Thr Trp Asp Lys Asn Val Arg Arg Gly Asp Ile Thr Ile Lys Leu Arg
            340
                                345
Asp Lys Ala Gly Asn Thr His Arg Ser Lys Ile Ile Ser Asn Glu Pro
                            360
                                                365
Thr Thr Phe Gln Lys Tyr His Gln Val Ser Leu Leu Ala Arg Phe Asn
                        375
Gln Asp Leu Asp Lys Val Ala Ala Ile Ser Leu Met Phe Ser Thr Gly
                    390
                                        395
Ser Leu Ile Gly Pro Arg Tyr Lys Leu Arg Ile Leu Arg Met Lys Leu
               405
                                    410
Arg Ser Leu Ala His Pro Glu Arg Pro Gln Leu Cys Arg Tyr Asp Leu
                                425
                                                    430
Val Leu Met Glu Asn Val Glu Thr Val Phe Gln Pro Ile Leu Cys Pro
                            440
       435
Glu Leu Gln Leu
    450
```

<211> 450

<212> PRT

<213> Homo sapiens

<400> 38

 Met
 Gly
 Leu
 Arg
 Ser
 His
 His
 Leu
 Ser
 Leu
 Gly
 Leu
 Leu
 Leu
 Phe

 Leu
 Pro
 Ala
 Glu
 Cys
 Leu
 Gly
 Ala
 Gly
 Arg
 Leu
 Ala
 Leu
 Leu
 Lys
 30
 Leu
 Lys
 30
 Leu
 Lys
 30
 Leu
 Lys
 30
 Lys
 45
 Lys
 <td

```
Ala Tyr Gly Gly Leu Asp Ala Ile Arg Ile Pro Ser Ser Leu Val Trp
          100
                       105
Arg Pro Asp Ile Val Leu Tyr Asn Lys Ala Asp Ala Gln Pro Pro Gly
                          120
                                              125
Ser Ala Ser Thr Asn Val Val Leu Arg His Asp Gly Ala Val Arg Trp
                       135
                                           140
Asp Ala Pro Ala Ile Thr Arg Ser Ser Cys Arg Val Asp Val Ala Ala
                   150
                                       155
Phe Pro Phe Asp Ala Gln His Cys Gly Leu Thr Phe Gly Ser Trp Thr
               165
                                   170
                                                       175
His Gly Gly His Gln Leu Asp Val Arg Pro Arg Gly Ala Ala Ala Ser
           180
                               185
Leu Ala Asp Phe Val Glu Asn Val Glu Trp Arg Val Leu Gly Met Pro
                           200
Ala Arg Arg Val Leu Thr Tyr Gly Cys Cys Ser Glu Pro Tyr Pro
                       215
                                            220
Asp Val Thr Phe Thr Leu Leu Leu Arg Arg Ala Ala Ala Tyr Val
                   230
                                       235
Cys Asn Leu Leu Pro Cys Val Leu Ile Ser Leu Leu Ala Pro Leu
                                   250
               245
Ala Phe His Leu Pro Ala Asp Ser Gly Glu Lys Val Ser Leu Gly Val
                               265
Thr Val Leu Leu Ala Leu Thr Val Phe Gln Leu Leu Ala Glu Ser
                           280
Met Pro Pro Ala Glu Ser Val Pro Leu Ile Gly Lys Tyr Tyr Met Ala
                       295
Thr Met Thr Met Val Thr Phe Ser Thr Ala Leu Thr Ile Leu Ile Met
                   310
                                       315
Asn Leu His Tyr Cys Gly Pro Ser Val Arg Pro Val Pro Ala Trp Ala
               325
                                   330
Arg Ala Leu Leu Gly His Leu Ala Arg Gly Leu Cys Val Arg Glu
                               345
           340
Arg Gly Glu Pro Cys Gly Gln Ser Arg Pro Pro Glu Leu Ser Pro Ser
                           360
Pro Gln Ser Pro Glu Gly Gly Ala Gly Pro Pro Ala Gly Pro Cys His
                       375
                                           380
Glu Pro Arg Cys Leu Cys Arg Gln Glu Ala Leu Leu His His Val Ala
                   390
                                       395
Thr Ile Ala Asn Thr Phe Arg Ser His Arg Ala Ala Gln Arg Cys His
                                   410
Glu Asp Trp Lys Arg Leu Ala Arg Val Met Asp Arg Phe Phe Leu Ala
           420
                               425
Ile Phe Phe Ser Met Ala Leu Val Met Ser Leu Leu Val Leu Val Gln
                           440
                                               445
Ala Leu
   450
<210> 39
<211> 255
<212> PRT
<213> Homo sapiens
Met Val Lys Gly Glu Lys Gly Pro Lys Gly Lys Lys Ile Thr Leu Lys
```

```
Val Ala Arg Asn Cys Ile Lys Ile Thr Phe Asp Gly Lys Lys Arg Leu
                               25
Asp Leu Ser Lys Met Gly Ile Thr Thr Phe Pro Lys Cys Ile Leu Arg
                           40
Leu Ser Asp Met Asp Glu Leu Asp Leu Ser Arg Asn Leu Ile Arg Lys
                      55
Ile Pro Asp Ser Ile Ser Lys Phe Gln Asn Leu Arg Trp Leu Asp Leu
                    70
                                       75
His Ser Asn Tyr Ile Asp Lys Leu Pro Glu Ser Ile Gly Gln Met Thr
                                    90
Ser Leu Leu Tyr Leu Asn Val Ser Asn Asn Arg Leu Thr Ser Asn Gly
           100
                                105
Leu Pro Val Glu Leu Lys Gln Leu Lys Asn Ile Arg Ala Val Asn Leu
                            120
Gly Leu Asn His Leu Asp Ser Val Pro Thr Thr Leu Gly Ala Leu Lys
                        135
Glu Leu His Glu Val Gly Leu His Asp Asn Leu Leu Asn Asn Ile Pro
                    150 -
                                        155
Val Ser Ile Ser Lys Leu Pro Lys Leu Lys Lys Leu Asn Ile Lys Arg
               165
                                    170
Asn Pro Phe Pro Lys Pro Gly Glu Ser Glu Ile Phe Ile Asp Ser Ile
           180
                                185
Arg Arg Leu Glu Asn Leu Tyr Val Val Glu Glu Lys Asp Leu Cys Ala
                           200
Ala Cys Leu Arg Lys Cys Gln Asn Ala Arg Asp Asn Leu Asn Arg Ile
                       215
Lys Asn Met Ala Thr Thr Pro Arg Lys Thr Ile Phe Pro Asn Leu
                   230
                                       235
Ile Ser Pro Asn Ser Met Ala Lys Asp Ser Trp Glu Asp Trp Arg
                                    250
```

<211> 214

<212> PRT

<213> Homo sapiens

<400> 40

Met Gln Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln Arg 10 Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln Pro 25 Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr Lys 40 Ile Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly Thr 55 60 Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln Glu 70 75 Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser Glu 90 Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Trp Glu Thr Lys Ile Asp 105 Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val Ile 120 125 Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn Val

```
Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile Asn
                        155
    150
Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg Ala
                              170
            165
Val Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val Ala
                          185
Glu Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu Glu
                         200
   195
Arg Cys Val Glu Ile Pro
   210
<210> 41
<211> 231
<212> PRT
<213> Homo sapiens
<400> 41
Met Met Pro Lys His Cys Phe Leu Gly Phe Leu Ile Ser Phe Phe Leu
                                 10
Thr Gly Val Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln
           20
                              25
                                                 30
Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln
                         40
                                            45
Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr
                     55
Lys Ile Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly
                 70
                                     75
Thr Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln
              85
                                 90
Glu Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser
          100
                             105
Glu Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Trp Glu Thr Lys Ile
                         120
                                            125
Asp Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val
   130 135
                                        140
Ile Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn
                 150
                                     155
Val Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile
              165
                                 170
                                                    175
Asn Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg
           180
                              185
                                                190
Ala Val Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val
                      200
                                             205
Ala Glu Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu
 210
                     215
                                         220
Glu Arg Cys Val Glu Ile Pro
225
                  230
<210> 42
<211> 263
<212> PRT
<213> Homo sapiens
```

<400> 42

```
Met Met Pro Lys His Cys Phe Leu Gly Phe Leu Ile Ser Phe Phe Leu
                                  10
Thr Gly Val Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln
                               25
Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln
                           40
                                                45
Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr
                       55
                                            60
Lys Ile Met Phe Ser Cys Ser Met Lys Ser Ser His Gln Lys Pro Ser
                   70
                                        75
Gly Cys Trp Gln His Ile Ser Cys Asn Phe Pro Gly Cys Arg Thr Leu
Ala Lys Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly
                                105
Thr Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln
                            120
Glu Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser
                        135
Glu Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Glu Thr Lys Ile
                   150
                                        155
Asp Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val
               165
                                    170
Ile Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn
                               185
Val Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile
                            200
Asn Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg
                       215
                                            220
Ala Val Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val
                   230
                                       235
Ala Glu Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu
                                    250
               245
Glu Arg Cys Val Glu Ile Pro
           260
```

<211> 259

<212> PRT

<213> Homo sapiens

<400> 43

Met Tyr Val Leu Ser Pro Val Glu Phe Ile Ile Leu Gln Leu Leu Phe 10 Ile Gln Ala Ile Ser Ser Ser Leu Lys Gly Phe Leu Ser Ala Met Arg 20 25 Leu Ala His Arg Gly Cys Asn Val Asp Thr Pro Val Ser Thr Leu Thr 40 45 Pro Val Lys Thr Ser Glu Phe Glu Asn Phe Lys Thr Lys Met Val Ile 55 60 Thr Ser Lys Lys Asp Tyr Pro Leu Ser Lys Asn Phe Pro Tyr Ser Leu 70 7.5 Glu His Leu Gln Thr Ser Tyr Cys Gly Leu Val Arg Val Asp Met Arg 90 Met Leu Cys Leu Lys Ser Leu Arg Lys Leu Asp Leu Ser His Asn His 105

```
Ile Lys Lys Leu Pro Ala Thr Ile Gly Asp Leu Ile His Leu Gln Glu
                            120
        115
Leu Asn Leu Asn Asp Asn His Leu Glu Ser Phe Ser Val Ala Leu Cys
                       135
                                            140
His Ser Thr Leu Gln Lys Ser Leu Arg Ser Leu Asp Leu Ser Lys Asn
                   150
                                        155
Lys Ile Lys Ala Leu Pro Val Gln Phe Cys Gln Leu Gln Glu Leu Lys
                                    170
                165
Asn Leu Lys Leu Asp Asp Asn Glu Leu Ile Gln Phe Pro Cys Lys Ile
                                185
                                                    190
Gly Gln Leu Ile Asn Leu Arg Phe Leu Ser Ala Ala Arg Asn Lys Leu
                            200
Pro Phe Leu Pro Ser Glu Phe Arg Asn Leu Ser Leu Glu Tyr Leu Asp
                        215
                                             220
Leu Phe Gly Asn Thr Phe Glu Gln Pro Lys Val Leu Pro Val Ile Lys
                    230
                                        235
Leu Gln Ala Pro Leu Thr Leu Leu Glu Ser Ser Ala Arg Thr Ile Leu
                245
                                    250
His Asn Arg
```

<210> 44 <211> 416 <212> PRT

<213> Homo sapiens

<400> 44

Met Lys Leu His Cys Glu Val Glu Val Ile Ser Arg His Leu Pro Ala 10 Leu Gly Leu Arg Asn Arg Gly Lys Gly Val Arg Ala Val Leu Ser Leu 20 25 Cys Gln Gln Thr Ser Arg Ser Gln Pro Pro Val Arg Ala Phe Leu Leu 40 45 Ile Ser Thr Leu Lys Asp Lys Arg Gly Thr Arg Tyr Glu Leu Arg Glu 5.5 Asn Ile Glu Gln Phe Phe Thr Lys Phe Val Asp Glu Gly Lys Ala Thr 75 Val Arg Leu Lys Glu Pro Pro Val Asp Ile Cys Leu Ser Lys Ala Ile Ser Ser Leu Lys Gly Phe Leu Ser Ala Met Arg Leu Ala His Arg 100 105 Gly Cys Asn Val Asp Thr Pro Val Ser Thr Leu Thr Pro Val Lys Thr 120 125 Ser Glu Phe Glu Asn Phe Lys Thr Lys Met Val Ile Thr Ser Lys Lys 135 140 Asp Tyr Pro Leu Ser Lys Asn Phe Pro Tyr Ser Leu Glu His Leu Gln 150 155 Thr Ser Tyr Cys Gly Leu Val Arg Val Asp Met Arg Met Leu Cys Leu 170 165 175 Lys Ser Leu Arg Lys Leu Asp Leu Ser His Asn His Ile Lys Lys Leu 185 190 Pro Ala Thr Ile Gly Asp Leu Ile His Leu Gln Glu Leu Asn Leu Asn 200 205 Asp Asn His Leu Glu Ser Phe Ser Val Ala Leu Cys His Ser Thr Leu

```
Gln Lys Ser Leu Arg Ser Leu Asp Leu Ser Lys Asn Lys Ile Lys Ala
225
                    230
                                         235
Leu Pro Val Gln Phe Cys Gln Leu Gln Glu Leu Lys Asn Leu Lys Leu
                                     250
                245
                                                          255
Asp Asp Asn Glu Leu Ile Gln Phe Pro Cys Lys Ile Gly Gln Leu Ile
            260
                                 265
                                                      270
Asn Leu Arg Phe Leu Ser Ala Ala Arg Asn Lys Leu Pro Phe Leu Pro
        275
                             280
                                                 285
Ser Glu Phe Arg Asn Leu Ser Leu Glu Tyr Leu Asp Leu Phe Gly Asn
    290
                        295
                                             300
Thr Phe Glu Gln Pro Lys Val Leu Pro Val Ile Lys Leu Gln Ala Pro
                    310
                                         315
                                                              320
Leu Thr Leu Leu Glu Ser Ser Ala Arg Thr Ile Leu His Asn Arg Asn
                                     330
Arg Ile Pro Tyr Gly Ser His Ile Ile Pro Phe His Leu Cys Gln Asp
            340
                                 345
                                                      350
Leu Asp Thr Ala Lys Ile Cys Val Cys Gly Arg Phe Cys Leu Asn Ser
                             360
                                                 365
Phe Ile Gln Gly Thr Thr Thr Met Asn Leu His Ser Val Ala His Thr
    370
                        375
                                             380
Val Val Leu Val Asp Asn Leu Gly Gly Thr Glu Ala Pro Ile Ile Ser
                    390
                                         395
Tyr Phe Cys Ser Leu Gly Cys Tyr Val Asn Ser Ser Asp Met Leu Lys
                                     410
```

```
<210> 45
<211> 990
<212> DNA
<213> Mus musculus
```

<400> 45

atggtggtca gggccttcgt tttgctggcc ctctttgcag aagcctcagc gaaatcatgc 60 actocgaata aagcagatgt catcottgtg ttttgttato ccaagaccat catcactaaa 120 atcoccgagt gtccctatgg atgggaagta caccagctgg cactcggggg gctgtgttac 180 aacggggtcc atgaaggtgg ctattaccag tttgtcatcc ctgatctgtc acctaagaac 240 aagtcctact gcggaaccca gtcagagtac aagcccccca tctaccactt ctacagccac 300 atogtgtcca acgacagcac agtgatcgtg aagaaccagc ccgtcaacta ctccttctcc 360 tgcacctacc actccaccta cttggtgaac caggctgctt ttgaccagag agtggccact 420 gttcacgtca agaacgggag catgggcaca tttgaaagcc agttgtccct caacttctac 480 actaatgcca agttttccac caaaaaagaa gctcccttcg ttctggaaac gtccgaaatc 540 ggctcagatc tgtttgcggg agtagaagcc aaaggcctaa gcgttcggtt caaagtggtc 600 ttgaataget getgggeeae eeectegget gaetteatgt acceettaca gtggeagete 660 atcaataagg gctgccccac cgatgagaca gtcctcgtgc atgagaacgg caaagaccac 720 agggccactt tccaattcaa tgccttccgg ttccagaaca tccccaaact ttccaaggtt 780 tggttacact gtgagacgtt catctgcgac agtgagaagc tctcctgccc cgtgaactgt 840 gacaaacgga agcgcatgct acgtgaccag acaggaggtg tcctggttgt ggagttgtcc 900 ctgaggagca gggcattttc cggcctctgt gacttctcag atgttcttct tcacctcatc 960 990 ctgatgctgg ggacctgggc tgtgttgtag

```
<210> 46
<211> 329
<212> PRT
<213> Mus musculus
```

```
Met Val Val Arg Ala Phe Val Leu Leu Ala Leu Phe Ala Glu Ala Ser
                                   10
Ala Lys Ser Cys Thr Pro Asn Lys Ala Asp Val Ile Leu Val Phe Cys
                               25
Tyr Pro Lys Thr Ile Ile Thr Lys Ile Pro Glu Cys Pro Tyr Gly Trp
                           40
                                               45
Glu Val His Gln Leu Ala Leu Gly Gly Leu Cys Tyr Asn Gly Val His
                        55
Glu Gly Gly Tyr Tyr Gln Phe Val Ile Pro Asp Leu Ser Pro Lys Asn
                    70
                                        75
Lys Ser Tyr Cys Gly Thr Gln Ser Glu Tyr Lys Pro Pro Ile Tyr His
                                    90
Phe Tyr Ser His Ile Val Ser Asn Asp Ser Thr Val Ile Val Lys Asn
            100
                                105
Gln Pro Val Asn Tyr Ser Phe Ser Cys Thr Tyr His Ser Thr Tyr Leu
                            120
                                                125
Val Asn Gln Ala Ala Phe Asp Gln Arg Val Ala Thr Val His Val Lys
                       135
                                            140
Asn Gly Ser Met Gly Thr Phe Glu Ser Gln Leu Ser Leu Asn Phe Tyr
                   150
                                        155
Thr Asn Ala Lys Phe Ser Thr Lys Lys Glu Ala Pro Phe Val Leu Glu
                                    170
               165
Thr Ser Glu Ile Gly Ser Asp Leu Phe Ala Gly Val Glu Ala Lys Gly
                                185
Leu Ser Val Arg Phe Lys Val Val Leu Asn Ser Cys Trp Ala Thr Pro
                           200
Ser Ala Asp Phe Met Tyr Pro Leu Gln Trp Gln Leu Ile Asn Lys Gly
                       215
                                            220
Cys Pro Thr Asp Glu Thr Val Leu Val His Glu Asn Gly Lys Asp His
                  230
                                       235
Arg Ala Thr Phe Gln Phe Asn Ala Phe Arg Phe Gln Asn Ile Pro Lys
               245
                                   250
Leu Ser Lys Val Trp Leu His Cys Glu Thr Phe Ile Cys Asp Ser Glu
           260
                               265
Lys Leu Ser Cys Pro Val Asn Cys Asp Lys Arg Lys Arg Met Leu Arg
                           280
                                               285
Asp Gln Thr Gly Gly Val Leu Val Val Glu Leu Ser Leu Arg Ser Arg
                       295
                                            300
Ala Phe Ser Gly Leu Cys Asp Phe Ser Asp Val Leu Leu His Leu Ile
                   310
                                        315
Leu Met Leu Gly Thr Trp Ala Val Leu
                325
```